polypeptide

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Database :
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Maximum Match 100%
Listing first 75 summaries
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Perfect score:
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Maximum DB
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Gapop 10.0 , Gapext 0.5
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                   | SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1980.DAT:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Res

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No.	Score	Query	Query Match Length DB	BB	ID	Description
ш	64	100.0	12	22	AAB74367	Reactive oxygen sp
N	64	100.0	12	22	AAB74370	Reactive oxygen sp
w	64	100.0	20	21	AAB12462	Human albumin epit
4	64	100.0	113	22	AA011631	Human polypeptide
ທ	64	100.0	116	22	AA002642	Human polypeptide
0	64	100.0	119	22	AA011632	
7	64	100.0	123	22	AA004424	Human polypeptide
8	64	100.0	124	22	AA002630	Human polypeptide
9	64	100.0	133	4.	AAP30004	Sequence encoded b
10	64	100.0	133	22	AAU29925	Novel human secret

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AAR125 AAR125 AAR057 AAR062 AAP3008 AAP3018 AAP6006 AAP6009 AAR253 AAR253 AAR253 AAR253	AAO2011 AAW5984 AAW8489 AAW84899 AAW848990 AABB9900 AABB1339 AABE1313 AABE1213 AABE12113 AABE12240 AABE12240 AABE123384 ABG63384 ABG63384 AAB75752 AAB70767	

B lymphocyte stimu Human B Lymphocyte Mature form of human Sequence of human Sequence of human Human serum albumi HSA-liver cell grous. HSA-fibronectin cells and HSA-nuclear transl Fragment of plasmi Sequence encoded by the state of the sequence encoded by the sequence Human serum albumi
Sequence of human
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HSA. Pichia pasto
HSA. Pichia pasto Human albumin (HA)
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Mature human serum
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ALIGNMENTS

RESULT 1
AAB74367
ID AAB7

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AAB74367 standard; peptide; 12 AA.

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RESULT 2
AAB74370
ID AAB7
XX AAB7
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XX ROS;
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                                                                                                                                                                                                                                                                                                                                                                                                                  Metal binding peptide compounds prevent damage by reactive oxygen species in animal organs and tissues, useful for reperfusion, transplantation and treating e.g. ischemia, neurological and cardiovascular diseases.
                                                                                                                                                                                                                                                                                                      Sequence
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13-JUN-2000; 2000US-0211078.
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                                  Synthetic
                                                     neurodegerative
                                                                  ROS; reactive oxygen
                                                                                        Reactive oxygen species inhibitory peptide #6.
                                                                                                               02-JUL-2001
                                                                                                                                                          AAB74370 standard; peptide; 12
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             WO200125265-A1.
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WINKLER J V.
CROOK W M.
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CURTIS C G.
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RESULT 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Metal binding peptide compounds prevent damage by reactive oxygen species in animal organs and tissues, useful for reperfusion, transmission and treating e.g. ischemia, neurological and
         Hybridoma cells for preparation of IgG monoclonal antibody capable of rapid reaction with human albumin for diagnosis of diabetic nephropathy
                                                                                                                                                                                                                  Monoclonal antibody; hybridoma cell; immunoglobulin; IgG; fusion; human albumin; diabetic nephropathy; diagnosis.
                                                                                                                                                                                                                                                   Human albumin epitope peptide #1.
                                                                                                                                                                                                                                                                                                AAB12462;
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                                            WPI; 2000-433935/38
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        rapid reaction with
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AISIN SEIKI
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CURTIS C G.
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ilarity 100.0%;
Conservative C
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2000US-0211078
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KK.
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            The invention relates to human polynucleotides (AAI79941-AAI93841) and the encoded proteins (AA000010-AA013910) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polynucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopoiesis regulating activity, tissue growth factor activity, immunomodulatory activity and activiny inhibin activity and may be useful in the diagnosis and/or activity introduced the control of th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; cytokine; cell proliferation; cell differentiation; gene the vaccine; peptide therapy; stem cell growth factor; haematopoiesis; tissue growth factor; immunomodulatory; cancer; leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-FEB-2000; 2000US-0515126.
18-MAY-2000; 2000US-0577409.
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                                                                                                                                                                                                                                                                                                                                                                                              Isolated nucleic acids and polypeptides, useful for preventing diagnosing and treating e.g. leukaemia, inflammation and immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-FEB-2001; 2001WO-US04927.
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                                                                                                                                                                                                                                                                                                            Claim 20; SEQ ID NO 25523; 1399pp + Sequence Listing; English.
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                                                                                       The invention relates to human polynucleotides (AAI79941-AAI93841) and the encoded proteins (AA000010-AA013910) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polynucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopoiesis regulating activity, tissue growth factor activity, immunomodulatory activity and activity inhibin activity and may be useful in the diagnosis and/or treatment of cancer, leukaemia, nervous system disorders, arthritis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; cytokine; cell proliferation; cell differentiation; gene therapy; vaccine; peptide therapy; stem cell growth factor; haematopoiesis; tissue growth factor; immunomodulatory; cancer; leukaemia; nervous system disorders; arthritis; inflammation.
                                                                                                                                                                                                                                                            Claim 20; SEQ ID NO 16534; 1399pp + Sequence Listing; English.
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18-MAY-2000;
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                                              specification,
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                                                                               inflammation.
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)B; AAI82573.
                                             The sequence data for this patent did not form part of the printed fication, but was obtained in electronic format directly from WIPO
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2000US-0577409.
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                                                                                                                                                specification,
                                                                                                                  Sequence
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                                                                                                                                                           Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                           inflammation.
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18-MAY-2000; 2000US-0577409.
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30
                                                                        Local Similarity
                          1 DAHKSEVAHREK 12
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DAHKSEVAHRFK 41
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Pred. No. 0.00018;
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                                                                                                                                                                                                                                                                                                                                                                                       cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polymucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopoiesis regulating activity, tissue growth factor activity, immunomodulatory activity and activin/inhibin activity and may be useful in the diagnosis and/or
                                               AAO02630;
                                                                             AAO02630 standard; Protein;
                                                                                                                                                                                                                                                                                   Sequence
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18-MAY-2000; 2000US-0577409.
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                                                                                                                                                                                                                                                                                   123 AA;
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Pred. No. 0.00019;
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06-NOV-2001

(first entry)

Human polypeptide SEQ ID NO 16522.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tang YT,
                                                                                  Sequence encoded by recombinant DNA molecule pKT218(HSA/33-1 (Bg1II-EcoRI)-HSA/17-3(Bg1II-EcoRI)) human proserum albumin (except for the missing 36 nucleotide Bg1II-Bg1II deletion).
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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   Homo sapiens
                                   Burn therapy; blood; protein; kernicterus therapy.
                                                                                                                                                          25-APR-1992
                                                                                                                                                                                          AAP30004;
                                                                                                                                                                                                                          AAP30004 standard; Protein; 133 AA
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12; Conservative
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                                                     nitrogen fixation; food additive;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Prodn. of human serum albumin-like polypeptide(s) - from host
transformed with recombinant DNA molecule
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1983-796520/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gilbert W, Philipp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HARD ) HARVARD COLLEGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-DEC-1981;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-MAY-1983;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EP91527-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          phase) in pKT218 still produce HSA-like products.
                                                                                                                                                                                                                      Human; vaccination; gene therapy; nutritional supplement; stem cell proliferation; haematopoiesis; nerve tissue regeneration; immune suppression; immune stimulation; anti-inflammatory; leukaemi
                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAU29925;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAU29925 standard; Protein; 133 AA
                                                                                                                  WO200179449-A2
                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                         Novel human secreted protein #416.
                                                                                                                                                                                                                                                                                                                                                                                                 18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27 DAHKSEVAHRFK 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 DAHKSEVAHRFK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     l Similarity
12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAN3 0009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     133 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          81US-0330912
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     83EP-0200712
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27..59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /label= residues 572-580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /label= residues 319-351
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /label= residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 64; DB 4;
Pred. No. 0.00021;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  368-399
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16-APR-2001; 2001WO-US08656

25-OCT-2001.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AA012088
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cc polypeptides and antibodies to the polypeptides are useful for cdetermining the presence of or predisposition to a disease associated with altered levels of polypeptide. The polypeptides are also useful for cdentifying agents (agonists and antagonists) that bind to them. Cells for use in treatment of a pathology related to aberrant expression or physiological interactions of the polypeptides and cells genetically cempineered to express them are also useful for producing the proteins. The proteins are useful in genetic vaccination, testing and in the therapy, and can be used as nutritional supplements. They may be used to therapy, and can be used as nutritional supplements. They may be used to cartilage, tendon and/or nerve tissue growth or regeneration; cartilage, tendon and/or nerve tissue growth or regeneration; commune suppression and/or stimulation; as anti-inflammacory agents; and contraction to regulate haematopolesis; and contraction to reduce anti-inflammacory agents; and contraction of leukaemias. AMU29510-AAU33304 represent the amino acid sequences of novel human secreted proteins of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Tang YT,
                                                                                      28-FEB-2000; 2000US-0515126.
18-MAY-2000; 2000US-0577409.
                                                                                                                                                      26-FEB-2001; 2001WO-US04927
                                                                                                                                                                                                                                               WO200164835-A2
                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                             Human; cytokine; cell proliferation; cell differentiation; gene therapy; vaccine; peptide therapy; stem cell growth factor; haematopoiesis; tissue growth factor; immunomodulatory; cancer; leukaemia; nervous system disorders; arthritis; inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Human polypeptide SEQ ID NO 25980.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AA012088;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAO12088 standard; Protein; 137 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acids encoding a range of human polypeptides, useful in genetic vaccination, testing and therapy - \,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-APR-2000; 2000US-0552929.
26-JAN-2001; 2001US-0770160.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-611725/70.
                                             (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to novel human secreted polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27 DAHKSEVAHREK 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 DAHKSEVAHRFK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12;
  Liu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 210; 765pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Liu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              133 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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  ņ
    Drmanac
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Pred. No. 0.00021;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 22; Length 133;
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AAU29875
ID AAU29
XX AAU29875
AC AAU29
AC AAU29
XX 18-DE
XX Human
KW Stem
KW Stem
KW Stem
KW Stem
KW 10000
XX W0200
XX 16-AP
PR 26-JA
XX 18-AP
PR 26-JA
XX IA-AP
PR 26-JA
XX IA-AP
PR 16-AP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                         Claim 20; Page 206; 765pp; English.
                                                                                                                                                      WPI; 2001-611725/70.
                                                                                                                                                                                                                                                                                                                                                                                                 25-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200179449-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                             (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inflammation.
invention relates to
                                                                                                                                                                                                 YT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                               Liu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       137 AA;
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the encoded proteins (AAOOOO10-AAO13910) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polymucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopoiesis regulating activity, tissue growth factor activity, hamonomodulatory activity and activity and may be useful in the diagnosis and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; vaccination; gene therapy; nutritional supplement; stem cell proliferation; haematopoiesis; nerve tissue regeneration; immune suppression; immune stimulation; anti-inflammatory; leukaemi
Nucleic acids encoding a range of human polypeptides, useful in genetic vaccination, testing and therapy -
                                                                                                                                                                                                                                                                                                                             18-APR-2000; 2000US-0552929.
26-JAN-2001; 2001US-0770160.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel human secreted protein #366.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAU29875 standard; Protein; 192 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     treatment of cancer, leukaemia, nervous system disorders, arthritis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 20; SEQ ID NO 25980; 1399pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated nucleic acids and polypeptides, useful for preventing diagnosing and treating e.g. leukaemia, inflammation and immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-514838/56.
N-PSDB; AAI92019.
                                                                                                                                                                                                                                                                                                                                                                                                                                 16-APR-2001; 2001WO-US08656
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30 DAHKSEVAHRFK 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 DAHKSEVAHRFK 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                     Drmanac RT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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Pred. No. 0.00021;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 137;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       leukaemia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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novel human secreted polypeptides.

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polypeptides and antibodies to the polypeptides are useful for determining the presence of or predisposition to a disease associated with altered levels of polypeptide. The polypeptides are also useful for identifying agents (agonists and antagonists) that bind to them. Cells expressing the proteins are useful for identifying a therapeutic agent for use in treatment of a pathology related to aberrant expression or physiological interactions of the polypeptide. Vectors comprising the nucleic acids encoding the polypeptides and cells genetically engineered to express them are also useful for producing the proteins. The proteins are useful in genetic vaccination, testing and therapy, and can be used as nutritional supplements. They may be used to increase stem cell proliferation; to regulate haematopoissis, and in the cartilage, tendom and/or nerve tissue growth or regeneration; immune suppression and/or stimulation; as anti-inflammatory agents; and in treatment of leukaemias. AAU29510-AAU33304 represent the amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
WPI; 2002-195801/25.
                                                 Hedley ML,
                                                                                                                                                          14-JUL-2000; 2000US-218381P.
18-AUG-2000; 2000US-236382P.
06-OCT-2000; 2000US-238380P.
29-DEC-2000; 2000US-258764P.
14-JUN-2001; 2001US-298317P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antiasthmatic; antibacterial; dermatological; antipsoriatic;
antidiabetic; ophthalmological; neuroprotective; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Alpha-MSH; inflammation; autoimmune disease; gene therapy; sepsis; alpha-melanocyte stimulating hormone; rheumatoid arthritis; asthma; cirrhosis; dermatitis; psoriasis; inflammatory bowel disease; immunosuppressive; antiinflammatory; antirheumatic; antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ery Match
                                                                                                                                                                                                                                                                                                                                  16-JUL-2001; 2001WO-US22263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200206316-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diabetes; uveitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human serum albumin (1-195)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAO17048;
                                                                                                        (ZYCO-) ZYCOS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 DAHKSEVAHRFK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HKSEVAHRFK 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of novel human secreted proteins of the invention.
                                                 Urban R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         192 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            coeliac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                 Aziz N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 64; DB
Pred. No. 0.0
0; Mismatches
                                                    Chen
                                                 Ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 22;
0.00032;
                                              Etemad-Moghadam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 192;
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                                                 Yin
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ક 밁

The present invention relates to a nucleic acid comprising a sequence

Query Match Best Local Similarity

100.0%;

Score 64; Pred. No.

DB 21; 0.00034;

Length 204;

Example 2; Page 46; 89pp; English.

disorders

Novel nucleic acid encoding fusion protein comprising alpha-melanocyte stimulating hormone concatamer or its analog, for treating inflammators

inflammatory

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RESULT 14
AAY83947
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                           The method relates to a method of recombinantly producing human serum albumin (HSA) in yeast by altering the coding sequence of HSA to comprise a yeast codon bias. The complete HSA gene (APA1091) was generated as three synthetic fragments (APA10092-A10094) joined by recombinant DNA technology. Each HSA fragment was synthesised from overlapping oligonuclectide fragments that were extended. This sequence represents the sequence of the HSA fragment HSA-I encoded by the human gene with a yeast codon bias. The invention also covers a recombinant expression vector, yeast host cells carrying the recombinant expression vector, yeast host cells carrying the recombinant expression vector and the process for producing human serum albumin in the yeast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        encoding a fusion polypeptide having an alpha-melanocyte stimulating hormone (MSH) concatamer. The sequences are useful for treating an individual suffering from, or at risk of, a disorder of the immune system e.g. inflammatory disorder or autoimmune disorder, including rheumatoid arthritis, asthma, sepsis, cirrhosis, dermatitis, pooriasis, contact hypersensitivity, inflammatory bowel disease, autoimmune encephalitis, multiple sclerosis, diabetes, lupus, uveitis and coeliac disease. The present sequence is a protein described in the exemplification of the
Sequence
                                host cell,
                                                                                                                                                                                                                            Example 1; Fig 3; 44pp; Chinese.
                                                                                                                                                                                                                                                                Process for preparing recombined human serum albumin - which comprises yeast biased sex codons
                                                                                                                                                                                                                                                                                                                N-PSDB; AAA10092
                                                                                                                                                                                                                                                                                                                                   Ψď.
                                                                                                                                                                                                                                                                                                                                                                Li S,
                                                                                                                                                                                                                                                                                                                                                                                                                                17-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Recombinant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yeast codon-biased recombinant HSA protein fragment HSA-I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY83947;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                               (HAIJ-) HAIJI BIOENGINEERING CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-DEC-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CN1239103-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              overlapping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAY83947 standard; Protein; 204 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 DAHKSEVAHRFK 12
                                                                                                                                                                                                                                                                                                                              2000-351198/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DAHKSEVAHRFK 12
                                                                                                                                                                                                                                                                                                                                                                Lu D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               l Similarity
12; Conserv
                             especially in secretory mode.
 204 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  195 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oligonucleotide; expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human serum albumin; HSA; yeast codon bias; host cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                98CN-0102506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98CN-0102506
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100.0%; F
ative 0;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 vector
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.00032;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 195;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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RESULT 15
AAU29874
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                                                                                                                                                                                                                                                                                                                                                    RSSSSSSSSSSSSSSS
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                            á
                                                                                                                                                   The invention relates to novel human secreted polypeptides. The CC polypeptides and antibodies to the polypeptides are useful for Cd determining the presence of or predisposition to a disease associated CC with altered levels of polypeptide. The polypeptides are also useful for cidentifying agents (agonists and antagonists) that bind to them. Cells expressing the proteins are useful for identifying a therapeutic agent for use in treatment of a pathology related to aberrant expression or physiological interactions of the polypeptide. Vectors comprising CC the nucleic acids encoding the polypeptides and cells genetically CC engineered to express them are also useful for producing the proteins. CC The proteins are useful in genetic vaccination, testing and CC increase stem cell proliferation; to regulate haematopoiesis; and in CC bone, cartilage, tendon and/or nerve tissue growth or regeneration; CC immune suppression and/or stimulation; as anti-inflammatory agents; and CC in treatment of leukaemias. AAU29510-AAU33304 represent the amino acid CC sequences of novel human secreted proteins of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                             Matches
                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; vaccination; gene therapy; nutritional supplement; stem cell prolkferation; haematopoiesis; nerve tissue regeneration; immune suppression; immune stimulation; anti-inflammatory; leukaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel human secreted protein #365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAU29874;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAU29874 standard; Protein; 214 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acids encoding a range of human polypeptides, useful in genetic vaccination, testing and therapy \boldsymbol{\cdot}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-APR-2000; 2000US-0552929.
26-JAN-2001; 2001US-0770160.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-APR-2001; 2001WO-US08656.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (HYSE-) HYSEQ INC.
                                                                                                                         Sequence
37
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                              1 DAHKSEVAHREK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DAHKSEVAHRFK 19
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DAHKSEVAHRFK 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12;
                                                            1 Similarity
12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 205;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Liu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                           214 AA;
                                                            100.0%;
ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    765pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                               0
                                                                              Score 64;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                               Mismatches
                                                               . 0.00036;
                                                                                               DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                             Length 214;
                                                                  Indels
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RESULT 16

AAO16984; 29-MAY-2002

(first entry)

AAO16984 standard; Protein; 241

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RESULT 17
AAO16984
                                                           В
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                                                                                                                    Matches
                                                                                                                                                Query Match
                                                                                                                                     Best
                                                                                                                                                                                                                     The present invention relates to a nucleic acid comprising a sequence encoding a fusion polypeptide having an alpha-melanocyte stimulating hormone (MSH) concatamer. The sequences are useful for treating an individual suffering from, or at risk of, a disorder of the immune system e.g. inflammatory disorder or autoimmune disorder, including rheumatoid arthritis, asthma, sepsis, cirrhosis, dermatitis, psoriasis, contact hypersensitivity, inflammatory bowel disease, autoimmune encephalitis, multiple sclerosis, diabetes, lupus, uveitis and coeliac disease. The present sequence is a peptide described in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-JUL-2000; 2000US-218381P.
18-AUG-2000; 2000US-286382P.
06-OCT-2000; 2000US-238380P.
29-DEC-2000; 2000US-258764P.
14-JUN-2001; 2001US-298317P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cirrhosis; dermatitis; psoriasis; inflammatory bowel disease; immunosuppressive; antiinflammatory; antirheumatic; antiarthritic; antiasthmatic; antibacterial; dermatological; antipsoriatic; antidiabetic; ophthalmological; neuroprotective; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human albumin-thrombin-alpha-MSH SEQ ID NO: 82.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA017051 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-JAN-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200206316-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diabetes; uveitis; coeliac disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           alpha-melanocyte stimulating hormone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Alpha-MSH; inflammation; autoimmune disease; gene therapy; sepsis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                       Novel nucleic acid encoding fusion protein comprising alpha-melanocyte stimulating hormone concatamer or its analog, for treating inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-JUL-2001; 2001WO-US22263.
                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                         or autoimmune disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-195801/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hedley ML,
                                                                                                                                                                                                                                                                                                                                                                             Example 2; Page 48; 89pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ZYCO-) ZYCOS INC.
                                                                                                                                                                                                             invention
                                                            25
                                                                                                                                     Local
                                                                                        1 DAHKSEVAHRFK 12
                                                            DAHKSEVAHRFK
                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Urban R,
                                                                                                                                                                               236 AA;
                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                     100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Aziz N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 236 AA
                                                                                                                       0
                                                                                                                                    Score 64; DB 23;
Pred. No. 0.0004;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chen H,
                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 rheumatoid arthritis; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Etemad-Moghadam
                                                                                                                                                     DB 23;
                                                                                                                                                     Length 236;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     B
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                                                                                                                          Gaps
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AAO16985
XX
XX
AC AAO1
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Z9-M
DT 29-M
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XX
XX
DB Alph
XX
XX
Alph
XW anti
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                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 18
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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-AUG-2000; 2
06-OCT-2000; 2
29-DEC-2000; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a nucleic acid comprising a sequence encoding a fusion polypeptide having an alpha-melanocyte stimulating hormone (MSH) concatamer. The sequences are useful for treating an individual suffering from, or at risk of, a disorder of the immune system e.g. inflammatory disorder or autoimmune disorder, including rheumatoid arthritis, asthma, sepsis, cirrhosis, dermatitis, psoriasis, contact hypersensitivity, inflammatory bowel disease, autoimmune encephalitis, multiple sclerosis, diabetes, lupus, uveits and coeliac disease. The present sequence is a protein described in the exemplification of the
             Alpha-MSH; inflammation; autoimmune disease; gene therapy; sepsis; alpha-melanocyte stimulating hormone; rheumatoid arthritis; asthma; cirrhosis; dermatitis; psoriasis; inflammatory bowel disease; immunosuppressive; antinflammatory; antirheumatic; antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alpha-MSH; inflammation; autoimmune disease; gene therapy; sepsis; alpha-melanocyte stimulating hormone; rheumatoid arthritis; asthma; cirrhosis; dermatitis; psoriasis; inflammatory bowel disease; immunosuppressive, antiinflammatory; antirheumatic; antiarthritic; antiasthmatic; antibacterial; dermatological; antipsoriatic; antiasthmatic; ophthalmological; neuroprotective; multiple sclerosis; antidiabetic; ophthalmological; neuroprotective; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel nucleic acid encoding fusion protein comprising alpha-melanocyte stimulating hormone concatamer or its analog, for treating inflammator:
   antiasthmatic;
                                                                                                 Alpha-MSH
                                                                                                                                                                                             AAO16985 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 2; Page 4-5; 89pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hedley ML,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-JAN-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diabetes; uveitis; coeliac disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Alpha-MSH
                                                                                                                                                                                                                                                                                                                                                                                                                                       invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ZYCO-) ZYCOS INC.
                                                                                                                                                                                                                                                                          25
                                                                                                                                                                                                                                                                                                       1 DAHKSEVAHRFK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                autoimmune disorders
                                                                                                                                                                                                                                                                          DAHKSEVAHRFK 36
                                                                                                                                                                                                                                                                                                                                         12;
                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                               construct
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; 2000US-226382P.
; 2000US-238380P.
; 2000US-258764P.
; 2001US-298317P.
                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                             (first
                                                                                                                                                                                                                                                                                                                                                                                                     B
 antibacterial;
                                                                                                                                                                                             Protein;
                                                                                            protein
                                                                                                                             entry)
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                                                                                              fragment SEQ ID NO: 70
                                                                                                                                                                                             242
                                                                                                                                                                                                                                                                                                                                                      Score 64;
Pred. No.
dermatological; antipsoriatic;
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                     Alpha-MSH; inflammation; autoimmune disease; gene therapy; sepsis; alpha-melanocyte stimulating hormone; rheumatoid arthritis; asthma cirrhosis; dermatitis; psoriasis; ninflammatory bowel disease; immunosuppressive; antiinflammatory; antirheumatic; antiarthritic; antiasthmatic; antibacterial; dermatological; antipsoriatic; antidabetic; ophthalmological; neuroprotective; multiple sclerosiantidiabetic; ophthalmological; neuroprotective; multiple sclerosiantidi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a nucleic acid comprising a sequence encoding a fusion polypeptide having an alpha-melanocyte stimulating hormone (MSH) concatamer. The sequences are useful for treating an individual suffering from, or at risk of, a disorder of the immune system e.g. inflammatory disorder or autoimmune disorder, including rheumatoid arthritis, asthma, sepsis, cirritosis, dermatitis, psoriasis, contact hypersensitivity, inflammatory bowel disease, autoimmune encephalitis, multiple sclerosis, diabetes, lupus, uveitis and coeliac disease. The present sequence is a protein described in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel nucleic acid encoding fusion protein comprising alpha-melanocyte stimulating hormone concatamer or its analog, for treating inflammatory
                                                                                                                          Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAO16986;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              or autoimmune disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hedley ML, Urban R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-JUL-2000;
18-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-JAN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-OCT-2000;
29-DEC-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                uveitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           construct protein fragment SEQ ID NO: 71.
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2000US-226382P.
2000US-238380P.
2000US-258764P.
2001US-298317P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001WO-US22263.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chen H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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hes 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 23;
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RESULT 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              stimulating hormone concatamer or autoimmune disorders -
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29-DEC-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-JUL-2000;
                                                                                                                                                                immunosuppressive; antiinflammatory; antirheumatic; antiarthritic;
antiasthmatic; antibacterial; dermatological; antipsoriatic;
antidiabetic; ophthalmological; neuroprotective; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                     AAO16987 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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                                                                                                                                                                                                       alpha-melanocyte stimulating hormone; rheumatoid arthritis; a cirrhosis; dermatitis; psoriasis; inflammatory bowel disease;
06-OCT-2000;
            14-JUL-2000;
18-AUG-2000;
                                                                                                     WO200206316-A2
                                                                                                                                                       diabetes; uveitis;
                                                                                                                                                                                                                                                          Alpha-MSH construct protein fragment SEQ ID NO: 72.
                                                                                                                                                                                                                                                                                    29-MAY-2002
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                                                 16-JUL-2001; 2001WO-US22263
                                                                          24-JAN-2002
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); 2000US-226382P.
); 2000US-238380P.
); 2000US-258764P.
; 2001US-298317P.
                                                                                                                                                                                                                                   inflammation; autoimmune disease;
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Pred. No.
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0.00042;
hes 0;
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                                                                                                                                                                                                                                   gene therapy;
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RESULT 21
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Best Local
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                                                                                                                                                                                  14-JUL-2000;
18-AUG-2000;
06-OCT-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel nucleic acid encoding fusion protein comprising alpha-melanocyte stimulating hormone concatamer or its analog, for treating inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-195801/25
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14-JUN-2001; 2001US-298317P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Alpha-MSH; inflammation; autoimmune disease; gene therapy; sepsis; alpha-melanocyte stimulating hormone; rheumatoid arthritis; asthma; cirrhosis; dermatitis; psoriasis; inflammatory bowel disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antiasthmatic; antibacterial; dermatological; antipsoriatic;
antidiabetic; ophthalmological; neuroprotective; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Alpha-MSH
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14-JUN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        immunosuppressive; antiinflammatory; antirheumatic; antiarthritic;
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                                                                               (ZYCO-) ZYCOS INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 DAHKSEVAHREK 12
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12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           uveitis;
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                                                                                                                                 2000US-226382P.
2000US-238380P.
2000US-258764P.
2001US-298317P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 coeliac
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                                   Etemad-Moghadam
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RESULT 22
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                                                                                                                                                                                                                                                                                                                                                                                                      Alpha-MSH; inflammation; autoimmune disease; gene therapy; sepsis; alpha-melanocyte stimulating hormone; fremenatoid arthritis; asthma; cirrhosis; dermatitis; psoriasis; inflammatory bowel disease; immunosuppressive; antinflammatory; antirheumatic; antiarthritic; antiasthmatic; antibacterial; dermatological; antipsoriatic; antidiabetic; ophthalmological; neuroprotective; multiple sclerosis; antidiabetic; ophthalmological; neuroprotective; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a nucleic acid comprising a sequence encoding a fusion polypeptide having an alpha-melanocyte stimulating hormone (MSH) concatamer. The sequences are useful for treating an individual suffering from, or at risk of, a disorder of the immune system e.g. inflammatory disorder or autoimmune disorder, including rheumatoid arthritis, asthma, sepsis, cirribosis, dermatitis, psoriasis, contact hypersensitivity, inflammatory bowel disease, autoimmune encephalitis, multiple sclerosis, diabetes, lupus, uveitis and coeliac disease. The present sequence is a protein described in the exemplification of the
                                                                                                                                                                                                           18-AUG-2000;
06-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                Unidentified
                                                                                                                                                                                                                                                                                                                                                                                              diabetes; uveitis; coeliac disease.
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                                         Novel nucleic acid encoding fusion protein comprising alpha-melanocyte stimulating hormone concatamer or its analog, for treating inflammator:
                                                                                      WPI; 2002-195801/25.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Alpha-MSH construct protein fragment SEQ ID NO: 60.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel nucleic acid encoding fusion protein comprising alpha-melanocyte stimulating hormone concatamer or its analog, for treating inflammatory
                                                                                                                                                  (ZYCO-) ZYCOS INC
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Example 2; Page 5; 89pp; English

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         polypeptides and antibodies to the polypeptides are useful for determining the presence of or predisposition to a disease associated with altered levels of polypeptide. The polypeptides are also useful for identifying agents (agonists and antagonists) that bind to them. Cells expressing the proteins are useful for identifying a therapeutic agent for use in treatment of a pathology related to aberrant expression or physiological interactions of the polypeptide. Vectors comprising the nucleic acids encoding the polypeptides and cells genetically engineered to express them are also useful for producing the proteins are useful in genetic vaccination, testing and the proteins are useful in genetic vaccination. They may be used to increase stem cell proliferation, to regulate haematopolesis, and in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to a nucleic acid comprising a sequence encoding a fusion polypeptide having an alpha-melanocyte stimulating hormone (MSH) concatamer. The sequences are useful for treating an individual suffering from, or at risk of, a disorder of the immune system e.g. inflammatory disorder or autoimmune disorder, including rheumatoid arthritis, asthma, sepsis, cirrhosis, dermatitis, pooriasis, contact hypersensitivity, inflammatory bowel disease, autoimmune encephalitis, multiple sclerosis, diabetes, lupus, uveitis and coeliac disease. The present sequence is a protein described in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; vaccinathon; gene therapy; nutritional supplement; stem cell proliferation; haematopoiesis; nerve tissue regeneration;
                                                                                                                                                                                                                      The invention relates to novel human secreted polypeptides. The
                                                                                                                                                                                                                                                                                           Nucleic acids encoding a range of human polypeptides, useful in genetic vaccination, testing and therapy -
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26-JAN-2001; 2001US-0770160.
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                                                                                                                                                                                                                                                        Claim 20; Page 167; 765pp; English.
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RESULT 24
AAU29581
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                                                                            polypeptides and antibodies to the polypeptides are useful for determining the presence of or predisposition to a disease associated with altered levels of polypeptide. The polypeptides are also useful for identifying agents (agonists and antagonists) that bind to them. Cells expressing the proteins are useful for identifying a therapeutic agent for use in treatment of a pathology related to aberrant expression or physiological interactions of the polypeptide. Vectors comprising the nucleic acids encoding the polypeptides and cells genetically can be useful in genetic vectors comprising the proteins are useful in genetic or producing the proteins. They may be used to therapy, and can be used as nutritional supplements. They may be used to increase stem cell proliferation; to regulate haematopoiesis; and in concease stem cell proliferation; to regulate haematopoiesis; and in treatment of leukaemias. ANU29510-ANU3304 represent the amino acid concease of novel human secreted proteins of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acids encoding a range of human polypeptides, useful in genetic vaccination, testing and therapy - \,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; vaccination; gene therapy; nutritional supplement; stem cell proliferation; haematopolesis; nerve tissue regeneration;
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                                                                                                                                                                                                                                                                                                                                                    The invention relates to novel human secreted polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                  Claim 20; Page 168; 765pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-611725/70
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26-JAN-2001; 2001US-0770160.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-APR-2001; 2001WO-US08656.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200179449-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel human secreted protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAU29581 standard; Protein; 289 AA.
                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immune suppression; immune stimulation; anti-inflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HYSE-) HYSEQ INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          37 DAHKSEVAHREK 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 DAHKSEVAHRFK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        l Similarity
12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Liu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          289 AA;
                                                  289 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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100.0%;
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 Score 64;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 64;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
 ; DB 22;
. 0.0005;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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                 Length 289;
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RESULT 25
AAR14178
ID AAR14
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AAP90387
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X S X F X G X W X S X G X G X
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This sequence corresponds to amino acids 1 to 303 of mature human serum albumin. The fragment lacking the C-terminal sequence can form part of a fusion protein, for example with drugs. (This sequence is taken from the full-length HSA sequence in EP-330451).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human serum albumin fragment - where C-terminal of human serum albumin is lacking and which can be combined with various drugs {\rm d} {\rm 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 1; 23pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-DEC-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-SEP-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HSA; fusion protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human serum albumin lacking C-terminal fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR14178;
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                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                             Human serum albumin polypeptide; plasma expanders
                                                                                                                                                                                                                                                                          N-terminal of human serum albumin polypeptide.
                                                                                                                                                                                                                                                                                                                                       01-NOV-1989
                                                                                                                                                                                                                                                                                                                                                                                                       AAP90387;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAP90387 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1991-300976/41.
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                            28-JUN-1989
                                                                                         EP322094-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
les 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       _
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DAHKSEVAHRFK 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     303 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                       (first entry
                                                                                                                                                      (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   89JP-0344701
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    373 AA
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Pred. No. 0.00053;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 303;
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RESULT 27
AAP90389
ID AAP90
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Matches
                          Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 373 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 2; page 9; 20pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New N-terminal fragments of human serum albumin - esp. useful as blood plasma expanders.
                                                                Sequence
                                                                                 N-terminal fragment of human serum albumin used as plasma expander, or as substitutes for HSA or BSA, in tissue culture media.
                                                                                                            Claim 2; page 9; 20pp; English.
                                                                                                                            New N-terminal fragments of human serum albumin - esp. useful as blood plasma expanders.
                                                                                                                                                                          Ballance DJ,
                                                                                                                                                                                                             30-OCT-1987;
                                                                                                                                                                                                                                                    28-JUN-1989.
                                                                                                                                                                                                                                                                     EP322094-A.
                                                                                                                                                                                                                                                                                                       N-terminal human serum albumin polypeptide; plasma expanders
                                                                                                                                                                                                                                                                                                                        N-terminal human serum albumin polypeptide.
                                                                                                                                                                                                                                                                                                                                            01-NOV-1989 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                AAP90389 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N-terminal portion of human serum albumin. or as substitutes for HSA or BSA, in tissue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ballance DJ, Hinchliffe E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-OCT-1987;
                                                                                                                                                         WPI; 1989-186464/26.
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                                                                                                                                                                                                                                                                                                                                                                AAP90389;
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                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                            (DELT ) DELTA BIOTECH LTD.
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                                   / Match 100.0%; Score 64; DB 10; Length 388; Local Similarity 100.0%; Pred. No. 0.00071;
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          1 DAHKSEVAHREK 12
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                           12; Conservative
                                                                388 AA;
                                                                                                                                                                                                                                                                                      (Human).
                                                                                                                                                                          Hinchliffe E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    87GB-0025529
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100.0%; Pred. No. 0.00068;
rative 0; Mismatches 0;
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                                                                                                                                                                          Geisow MJ,
                                                                                                                                                                                                                                                                                                                                                                                 388 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Geisow MJ,
                            Mismatches
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                                                                                                                                                                           Senior PJ;
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ID AAP90
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PA (DEL1)
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                                    25-OCT-1988;
                                                                                                           N-terminal portion of human
                                                                                                                             N-terminal human serum albumin.
                                                                                                                                                                AAP90391;
                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                    Claim 2; page 9; 20pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                   Ballance DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-OCT-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EP322094-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N-terminal human serum albumin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-NOV-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAP90390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAP90390 standard; protein;
                  30-OCT-1987;
                                                      28-JUN-1989
                                                                        EP322094-A.
                                                                                         Homo sapiens (Human).
                                                                                                                                               01-NOV-1989
                                                                                                                                                                                  AAP90391 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1989-186464/26.
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                                                                                                                                                                                                                                                                                                        389 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                  Hinchliffe E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (Human).
                   87GB-0025529
                                     88EP-0310000.
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Query Match 100.0%; Score 64; DB 10; Length 389; Best Local Similarity 100.0%; Pred. No. 0.00071; Matches 12; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N-terminal portion of human serum albumin. Used to make plasma expanders, or as substitutes for HSA or BSA, in tissue culture media.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New N-terminal fragments of human serum albumin - esp. useful as blood plasma expanders.
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(DELT ) DELTA BIOTECH LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Senior PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                <u>.</u>.
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RESULT 30
AAU29876
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
The invention relates to novel human secreted polypeptides. The polypeptides and antibodies to the polypeptides are useful for determining the presence of or predisposition to a disease associated with altered levels of polypeptide. The polypeptides are also useful for identifying agents (agonists and antagonists) that bind to them. Cells expressing the proteins are useful for identifying a therapeutic agent for use in treatment of a pathology related to aberrant expression or physiological interactions of the polypeptide. Vectors comprising the nucleic acids encoding the polypeptides and cells genetically engineered to express them are also useful for producing the proteins. The protesins are useful in genetic vaccination, testing and therapy, and can be used as nutritional supplements. They may be used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; vaccination; gene therapy; nutritional supplement; stem cell proliferation; haematopoiesis; nerve tissue regeneration; immune suppression; immune stimulation; anti-inflammatory; leukaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N-terminal portion of human serum albumin. Used to make n N-terminal fragments which are used as plasma expanders, or as substitutes for HSA or BSA, in tissue culture media.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                  Nucleic acids encoding a range of human polypeptides, useful in genetic vaccination, testing and therapy \mbox{-}
                                                                                                                                                                                                                                                                                   WPI; 2001-611725/70.
                                                                                                                                                                                                                                                                                                                                                                            18-APR-2000; 2000US-0552929.
26-JAN-2001; 2001US-0770160.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel human secreted protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAU29876;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAU29876 standard; Protein; 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 2; page 9; 20pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New N-terminal fragments of human serum albumin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1989-186464/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ballance DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                        16-APR-2001; 2001WO-US08656.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-OCT-2001
                                                                                                                                                                                                      Claim 20; Page 206; 765pp; English.
                                                                                                                                                                                                                                                                                                                                               (HYSE-) HYSEQ INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200179449-A2
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nes 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 DAHKSEVAHRFK 12
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                                                                                                                                                                                                                                                                                                                Liu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           390 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hinchliffe E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  blood plasma expanders
                                                                                                                                                                                                                                                                                                                Drmanac RT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 64; DB 10;
Pred. No. 0.00071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Senior PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Used to make new
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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RESULT 32 AAU29877

18-DEC-2001

(first entry)

AAU29877;

AAU29877 standard; Protein; 550 AA

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RESULT 31
AAP90392
ID AAP90
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                                                                                                          Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                             N-terminal portion of human serum albumin. plasma expanders, or as substitutes for HSA in tissue culture media.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         increase stem cell proliferation; to regulate haematopoiesis; and in bone, cartilage, tendon and/or nerve tissue growth or regeneration; immune suppression and/or stimulation; as anti-inflammatory agents; and in treatment of leukaemias. AAU29510-AAU33304 represent the amino acid sequences of novel human secreted proteins of the invention.
                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                      New N-terminal fragments of human serum albumin - esp. useful as blood plasma expanders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         plasma expanders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human serum albumin; mature protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N-terminal human serum albumin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-NOV-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAP90392;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                 Claim 2; page 9; 20pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1989-186464/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ballance DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-OCT-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-OCT-1988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (DELT ) DELTA BIOTECH LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              38 DAHKSEVAHRFK 49
\vdash
                                                  1 DAHKSEVAHRFK 12
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DAHKSEVAHRFK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; milarity 100.0%; Conservative 0;
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                                                                                                                                                                                                                           407 AA;
                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hinchliffe E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      87GB-0025529
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                                                                                                                                     100.0%; Score 64; DB 10;
100.0%; Pred. No. 0.00075;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Geisow MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 64;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       new
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. 0.00074;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Senior PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polypeptides;
                                                                                                                                                                                                                                                                                                              Used to make or BSA,
                                                                                                                                                                   Length 407;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
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                                                                                                                Gaps
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Novel human secreted protein #368.

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                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                            Sequence of mature human serum albumin (HSA) as
Homo sapiens
                                               Mature human serum albumin;
                                                                                                                              23-JUN-1990
                                                                                                                                                                                         AAP93344 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                hypobolaemia;
                                                                               artificial gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 20; Page 206; 765pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acids encoding a range of human polypeptides, useful in genetic vaccination, testing and therapy -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; vaccination; gene therapy; nutritional supplement; stem cell proliferation; haematopoiesis; nerve tissue regeneration; immune suppression; immune stimulation; anti-inflammatory; leukaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to novel human secreted polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-611725/70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-APR-2000; 2000US-0552929.
26-JAN-2001; 2001US-0770160.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HYSE-) HYSEQ INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-APR-2001; 2001WO-US08656.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200179449-A2
                                                                                                                                                                                                                                                                            27 DAHKSEVAHRFK 38
                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                           1 DAHKSEVAHRFK 12
                                                                                                                                                                                                                                                                                                                                         12;
                                                                                                                                                                                                                                                                                                                                                                                                         550 AA;
                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                           (first entry)
                                shock; hypoalbuminaemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Drmanac RT
                                                                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                                                                                       100.0%;
                                           artificial
                                                                                                                                                                                                                                                                                                                                         0,
                                                                                                                                                                                                                                                                                                                                                     Score 64; DB 22;
Pred. No. 0.0011;
                                                                                                                                                                                                                                                                                                                                         Mismatches
                                           gene; oligonucleotide block;
                                                                                                                                                                                                                                                                                                                                                                       DB 22;
                                                                                              encoded
                                                                                                                                                                                                                                                                                                                                                                    Length 550;
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RESULT 34
AAP90388
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The synthetic gene was constructed by designing a nucleotide sequence in which the codons which are most frequently used by the chosen non-human host were selected. In this case, it is yeast cells (LL20; Leu2-3; 112, His 3-11, 15). The synthetic HSA gene was assembled from 24 oligonucleotide blocks. HSA is used in therapy for the treatment of hypovolaemia, shock and hypoalbuminaemia. It is also used as an additive in perfusion liq. for extracorporeal circulation and as an experimental
                                                                                            30-OCT-1987;
                                                                                                                25-OCT-1988;
                                                                                                                                      28-JUN-1989
                                                                                                                                                                              Homo sapiens (Human)
                                                                                                                                                                                                                                                         01-NOV-1989
                  N-PSDB; AAN90128.
                                                 Ballance DJ,
                                                                       (DELT ) DELTA BIOTECH LTD.
                                                                                                                                                                                                    piasma expanders.
                                                                                                                                                                                                            Human serum albumin; mature protein; new polypeptides;
                                                                                                                                                                                                                                 Mature human serum albumin polypeptide.
                                                                                                                                                                                                                                                                              AAP90388;
                                                                                                                                                                                                                                                                                                 AAP90388 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; pp. 11-16; 121pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Artificial gene coding for authentic human serum albumin constructed of the basis of codons most frequently used by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-SEP-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chosen non-human host
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Aberg B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-SEP-1988;
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                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                       ب
                                                                                                                                                                                                                                                                                                                                                                          1 DAHKSEVAHRFK 12
                             1989-186464/26.
                                                                                                                                                                                                                                                                                                                                                       DAHKSEVAHRFK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1989-087749/12.
                                                                                                                                                                                                                                                                                                                                                                                                 12;
                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SKANDIGEN AB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAN90997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Simoncsits A,
                                                                                                                                                                                                                                                                                                                                                                                                                                          585 AA;
                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                         (first entry)
                                                 Hinchliffe E,
                                                                                           87GB-0025529
                                                                                                                88EP-0310000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       87SE-0003539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88EP-0850299
                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 64; DE 100.0%; Pred. No. 0.0 tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MAGY VEPE-).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kalman M,
                                                 Geisow
                                                                                                                                                                                                                                                                                                   585
                                                                                                                                                                                                                                                                                                   A
                                                 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cserpan I,
                                                                                                                                                                                                                                                                                                                                                                                                         0.0011;
                                                 Senior PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                   DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bajszar
                                                                                                                                                                                                                                                                                                                                                                                                                  Length 585;
                                                                                                                                                                                                                                                                                                                                                                                               Indels
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New N-terminal fragments of human

serum albumin

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RESULT 35
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Best Local
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                                                          Query Match
Best Local Similarity
                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mature protein of human serum albumin (see corresp. AAN90128). Used to make new N-terminal fragments which are used as plasma expanders, or as substitutes for HSA or BSA, in tissue culture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human normal serum albumin A; pAT-pho-HSA-A; haemorrhagic shock; hypoalbuminaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human normal serum albumin A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAP91422;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAP91422 standard; protein; 585 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; fig 2; 20pp; English.
                                                                                                                     cDNA amino acid sequence of human serum albumin A (HSA-A) which is identical to that encoded by chromosomal DNA. Previous polypeptides produced from cDNA have one or more amino acids which differ from those of HSA-A produced from the chromosomal DNA, and may exhibit antigenicity when administered to humans. The HSA-A is used to treat haemorrhagic shock and hypoalbuminaemia. See also AAN90600.
                                                                                                                                                                                                                                                                                                                                                                                                                                    EP330451-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-DEC-1989
                                                                                                                                                                                                                                                                                                                                                                                                             30-AUG-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                             cDNA encoding human normal serum albumin contained in plasmid - obtd. by
                                                                                                                                                                                                                                                                                                            Susuki M,
                                                                                                                                                                                                                                                                                                                                                            22-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                    22-FEB-1989;
                                                                                                Seguence
                                                                                                                                                                                                          Claim 1; fig3-1 to 3-5; 19pp; English.
                                                                                                                                                                                                                                                                                                                                   (TOFU ) TOA NENRYO KOGYO KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      esp. useful as blood plasma expanders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 DAHKSEVAHRFK 12
              1 DAHKSEVAHRFK 12
1 DAHKSEVAHREK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DAHKSEVAHRFK 12
                                                12;
                                                                                                                                                                                                                                                                                                            Maki N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            585 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 64; DB 10; ilarity 100.0%; Pred. No. 0.0011; Conservative 0; Mismatches 0
                                                                                                585 AA;
                                              100.0%; Score 64; DB 10; ilarity 100.0%; Pred. No. 0.0011; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                            88JP-0037453.
                                                                                                                                                                                                                                                                                                                                                                                    89EP-0301731.
                                                                                                                                                                                                                                                                                                            Yagi S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10;
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                                                                          Length 585;
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AAR05318
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XX JP02
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XX Graph (TO)
PA (TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 37
AAR08457
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         멍
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                    Query Match
Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human serum albumin; HSA-A; yeast; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human serum albumin gene product
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-OCT-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR05318;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mature HSA-A may be produced using the sequence incorporated into a plasmid vector with suitable controllers, and transfered to a yeast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; ; pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human serum albumin prepn. by yeast host -
by culturing transformed plasmid yeast to produce serum, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N-PSDB; AAQ04719.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1990-176228/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-OCT-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-OCT-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAY-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JP02117384-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        expression system.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       removing it.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR08457 standard; Protein; 585 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (TOFU ) TOA NENRYO KOGYO KK
                                                                                                                                                                                                                                                                                               HSA; folding; ss
                                                                                                                                                                                                                                                                                                                                                 Human serum albumin
                                                                                                                                                                                                                                                                                                                                                                                                  16-APR-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR08457;
JP02227079-A.
                                                                   Region
                                                                                                                       Region
                                                                                                                                                                        Region
                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
les 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 DAHKSEVAHRFK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 DAHKSEVAHRFK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               585 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         88JP-0268302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88JP-0268302
                                                                                                                       Location/Qualifiers
123..303
/label= A
1..303
                                              /label= B
123..585
/label= C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
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Pred. No. 0.0011;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 585;
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RESULT 38
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HSA can be recombinantly produced in yeast. A HSA coding sequence is incorporated into a vector which also contains a DNA region having the same base sequence as that of the target region of the chromosome and a promoter (esp. HIS4 or LEU2) to regulate the expression of the HSA coding sequence in the yeast host.
                                                                                                                          Stable prepn. of human serum albumin - by culturing yeast in which plasmid for recombining DNA coding human serum albumin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            fragment, C, has good star
segment, A, has character
See also AAQ06096-Q06098.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pragments A-C of HSA are expressed as fusion proteins with the signal peptide of E. coli alkaline phosphatase. The fragments are selected for their specific properties. The C-terminal truncated fragment, B, does not bind long-chain fatty acids but does bind to various medicines at the central region. The N-terminal truncated fragment, C, has good stability in protein folding. The central segment, A, has characteristics of both B and C.
                                                                                    Disclosure; Fig 4; 12pp; Japanese
                                                                                                                                                                         N-PSDB; AAQ20201.
                                                                                                                                                                                                                                                                                                                              JP03262487-A.
                                                                                                                                                                                                                                                                                                                                                                                                                  Human serum albumin
                                                                                                                   etc. is inserted
                                                                                                                                                                                                                                             12-MAR-1990;
                                                                                                                                                                                                                                                                        12-MAR-1990;
                                                                                                                                                                                                                                                                                                  22-NOV-1991
                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                     HSA; yeast promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                              08-APR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR20029 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New human serum albumin fragments - used to bond medicines and stable folding of protein(s).
                                                                                                                                                                                                                 (TOFU ) TONEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Fig 8; 24pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-OCT-1988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-SEP-1990;
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                                                                                                                                                                                      1992-012704/02
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.0011
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XX
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Best Local S
Matches 12
                             Query Match
Best Local 9
          Matches
                                                                                                        Sequence
                                                                                                                                                   The expression vector encoding this protein can be used for the large scale production of HSA. The HSA can then be administered to patient with circulatory failure or albumin depletion.
                                                                                                                                                                                                                                                      mating factor (AMF) pre-pro sequence (AAR26208) or the serum albumin (HSA) signal sequence; and
(ii) an HSA peptide (AAR26207); and
(c) a transcription terminator, functional in
                                                                                                                                                                                                                                                                                                                                                                                                               a methylotrophic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA fragment for prodn. of human serum albumin - comprises expression cassette including promoter and terminator sequences of methanol responsive gene, for expression in methylotrophic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  containing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The sequences given in AAR26207-9 were encoded by an expression cassette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 50-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1992-300037/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Davis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-FEB-1991;
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                                                                                                                                                                                                                                   Yeast (AAR26209).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               alpha-mating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human serum albumin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-FEB-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR26207 standard; Protein; 585 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Expression cassette;
alpha-mating factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (SALK ) SALK INST BIOTECHNOLOGY IND ASSOC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GR,
     12;
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                                Similarity
                                                                                                                                                                                                                                                                                                                                                      (b) a DNA sequence encoding a polypeptide encoding (i) a secretion signal from either S. cerevisa
                                                                                                                                                                                                                                                                                                                                                                                                                                    (a) the promoter region of a methanol-responsive
                                                                                                           585 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       585 AA;
     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%;
ilarity 100.0%;
Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                 yeast;
                        100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               methanol-responsive gene; methylotropic yeast; \mathsf{AMF}; \mathsf{HSA}.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       75pp; English.
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                          Score
Pred.
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Pred. No. 0.0011;
Pred. No. 0.0
Mismatches
                          0.0011
                                                     DB 13;
                                                                                                                                                                                                                                                      functional in methylotropic
                                                                                                                                                                           administered to patients
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 585;
     Indels
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                                                                                                                                                                                                                                                                                                                                 cerevisae alpha-
the native human
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Gaps
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RESULT 40
AAR26362
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                                                                                                                                                                                                                                                                                                       Query Match
Best Local (
                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JP04211375-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human serum albumin; transformants; recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic HSA protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-FEB-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAR26362 standard; Protein; 585 AA
                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic gene for prepn. of human serum albumin -synthetic DNA contg. gene coding the albumin using Escherichia coli
                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1992-304940/37.
N-PSDB; AAQ27813.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-FEB-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-FEB-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-AUG-1992.
                                                                                                                                                                                                                                                                                                                                                                The protein sequence was deduced from the synthetic DNA sequence encoding human serum albumin which was prepd. by ligating eight
                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                       synthetic HSA gene fragments.
                                                                                                                                                                                                                                                                                                                                                                                               Fig 1; Page 13; 37pp; Japanese.
                                                                                                          Saccharomyces cerevisiae.
                                                                                                                   Serum albumin; HSA; aspartyl protease-3; Yap3p
                                                                                                                                         Human serum albumin
                                                                                                                                                              17-JAN-1996 (first entry)
                                                                                                                                                                                                     AAR80301 standard; Protein; 585 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (AJIN ) AJINOMOTO KK.
                                                                   WO9523857-A1
                                                                                        Homo sapiens
          05-MAR-1994;
                             01-MAR-1995;
                                                 08-SEP-1995
                                                                                                                                                                                                                                                       _
                                                                                                                                                                                                                                                                          1 DAHKSEVAHREK 12
                                                                                                                                                                                                                                                       DAHKSEVAHREK 12
                                                                                                                                                                                                                                                                                              12;
                                                                                                                                                                                                                                                                                                      h 100.0%;
Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                     585 AA;
                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90JP-0025682
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                91JP-0014600
          94GB-0004270
                             95WO-GB00434.
                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                        Score 64; DB 13;
Pred. No. 0.0011;
                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                   DB 13;
                                                                                                                                                                                                                                                                                                                   Length 585;
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RESULT 42
AAO20111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The cDNA given in AAQ98695, which encodes HSA (AAR80301), was subjected to site-directed mutagenesis to investigate the role of endoproteases in the generation of a 45 kDa albumin fragment obtd. when the cDNA is expressed in S. cerevisiae. Mutations were: R410A; L407A, L408V, V499A; and R410A, K413Q, K414Q. The latter set of mutations, especially, improved stability of HSA to yeast Yap3P proteolytic cleavage, allowing increased prodn. of recombinant HSA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 26-28; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      kD fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yeast with reduced levels of aspartyl protease 3 proteolytic activity - used to secrete human albumin without prodn. of the 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1995-320572/41.
N-PSDB; AAQ98695.
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                                   The invention relates to a serum albumin-growth hormone fusion protein - useful to treat growth hormone related diseases such as Down's syndrome. This sequence represents a HSA protein related to the serum albumin-growth hormone protein of the invention.
                                                                                                                                                                   Serum albumin-growth hormone fusion protein - useful to treat growth hormone related diseases, e.g. Down's syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HSA protein sequence related to the growth hormone protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-AUG-2002
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                                                                                                                                                                                                                            N-PSDB; AAK99568.
                                                                                                                                                                                                                                                 WPI; 1997-363680/55.
                                                                                                                                                                                                                                                                                                                       30-DEC-1995;
19-DEC-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Down's syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Serum albumin-growth hormone fusion protein; growth hormone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAO20111;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAO20111 standard; Protein; 585 AA.
Sequence
                                                                                                                                Disclosure; Fig 6; 21pp; Korean.
                                                                                                                                                                                                                                                                                                                                                                                                                   15-0CT-1999
                                                                                                                                                                                                                                                                                  (DELZ ) DELTA BIOTECHNOLOGY LTD
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nes 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 DAHKSEVAHRFK 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kerry-Williams SM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                     95GB-0026733.
96WO-GB03164.
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Pred. No. 0.0011;
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Query Match

Local Similarity

100.0%; Score 64; 100.0%; Pred. No.

0.0011 DB 18;

Length 585;

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                                                                    RESULT 43
                                                                                             The present sequence represents the mature protein of human serum CC albumin (HSA). The protein is used to exemplify the invention. The specification describes a method for producing mature heterologous CC protein in monocotyledonous plant cells. The method comprises CC transforming the cells with a chimeric gene comprising a monocotyledon curanscription regulator, inducible either during seed maturation or by adding/removing a small molecule, DNA encoding the heterologous protein, CC and DNA encoding a signal peptide, with the signal peptide causing secretion of the protein from the cell. Proteins expressed in this canner include mature glycosylated alpha 1-antitrypsin (AAT) with a glycosylation pattern that significantly increases its serum half-life, CC mature glycosylated antithrombin III (ATIII), mature human serum albumin CC (HSA) having the native folding pattern as shown by bilirubin-binding CC characteristics, or mature active subtilisin BPN'. These proteins are CC useful therapeutically (e.g. AAT for treating emphysema, ATIII as CC antithrombotic and HSA as blood replacement) or as industrial enzymes CC (BPN' is used in detergents).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                            Expressing mature, glycosylated proteins in monocotyledonous plant cells - from chimeric gene including signal peptide sequence, specifically therapeutic agents and industrial enzymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Protein expression; monocotyledon plant cell; glycosylated antithrombin III; glycosylated alpha 1-antitrypsin; AAT; glycosylated antithrombin III; ATIII; human serum albumin; HSA; subtilisin BPN'; treatment; emphysema; antithrombotic; blood replacement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1998-467179/40.
N-PSDB; AAV41728.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Pages 31-32; 53pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rodriguez RL,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-FEB-1998;
                                                                    Seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (PHYT-) APPLIED PHYTOLOGICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-FEB-1997;
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                                                                    585 AA;
   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sutliff
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97US-0038169
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97US-0037991.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human serum albumin (HSA).
100.0%; S
100.0%; F
tive 0;
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                   Score 64; DB 19;
Pred. No. 0.0011;
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   Mismatches
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   0;
                                 Length 585;
 Indels
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RESULT 44
AAY84873
ID AAY84
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                                                            밁
                                                                                   The present sequence represents human albumin protein. The specification cd describes a method for the continuous detection of ischemic states. The method comprises detecting and quantifying the existence of an alteration of the serum protein albumin. The method comprises contacting a biological sample containing albumin from the patient with an excess quantity of a metal ion salt, where the metal ion binds to the N-terminus of naturally occurring human albumin, to form a mixture containing bound containing bound metal ions and unbound metal ions, and then determining the amount of metal ions bound to the albumin N-terminus. The amount of bound metal ions is correlated to a known value to determine the occurrence or non-occurrence of an ischemic event. The methods are useful for detection of ischemic states. The methods are also useful for distinguishing perloperative ischemia from ischemia caused by , amongst other things, compocardial infarctions and progressive coronary artery disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02-OCT-1998;
02-OCT-1998;
02-OCT-1998;
11-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                    New method for the continuous detection of ischemic states comprises detecting and quantifying the existence of an alteration of the seru
                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-303843/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bar-Or D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; albumin; ischemic state; serum protein; metal ion salt; perioperative ischemia; ischemia; myocardial infarction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Amino acid sequence of a human albumin protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAY84873;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY84873 standard; protein; 585 AA
                                                            Sequence
                                                                                                                                                                                                                                                                                                                         Disclosure; Page 97-100; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                      protein albumin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         progressive coronary artery disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISCH-) ISCHEMIA TECHNOLOGIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          μ
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              Similarity
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                                                            585
 Conservative
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98US-0165581.
98US-0165926.
99US-0115392.
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    claim 56"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Winkler JV;
              100.0%;
   0
Score 64; DB 21; Pred. No. 0.0011; Mismatches 0
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                               DB 21;
                              Length 585;
   Indels
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   Gaps
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1 DAHKSEVAHRFK 12

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RESULT 46
ABB79006
ID ABB79
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ABB79
XY
AC ABB79
DT 01-AU
XX
DE Human
XX
KW Human
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                                                                                                                                                                         Matches
                                                                                                                                                                                 Query Match
Best Local (
                                                                                                                                                                                                                                        The method relates to a method of recombinantly producing human serum albumin (HSA) in yeast by altering the coding sequence of HSA to comprise a yeast codon bias. The complete HSA gene (AAA10091) was generated as three synthetic fragments (AAA10092-A10094) joined by recombinant DNA technology. Each HSA fragment was synthesised from overlapping oligonucleotide fragments that were extended. This sequence represents the complete sequence of the HSA encoded by the human gene with a yeast codon bias. The invention also covers a recombinant expression vector, yeast host cells carrying the recombinant expression vector and the process for producing human serum albumin in the yeast host cells carry mode.
                   Human mature albumin protein SEQ ID NO:18.
                                                                                                                                                                                                                Sequence
 Human; growth hormone; hGH; albumin; human serum albumin; HSA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Recombinant; human serum albumin; HSA; yeast codon bias; host cell; overlapping oligonucleotide; expression vector.
                                        01-AUG-2002
                                                                               ABB79006 standard;
                                                                                                                                                                                                                                                                                                                                                       Disclosure; Fig 1; 44pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                           Process for preparing recombined human serum albumin - which comprises
yeast blased sex codons
                                                                                                                                                                                                                                                                                                                                                                                                                                       Li S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAY83946 standard; Protein; 585 AA.
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1 DAHKSEVAHRFK 12
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                                                                                                                                                                                  Similarity
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                                                                                                                                                                                                                585 AA;
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                                                                                                                                                                        Conservative
                                       (first entry)
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                                                                                Protein; 585
                                                                                                                                                                                 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     recombinant human serum albumin protein.
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                                                                                                                                                                                 Score 64; DB 21;
Pred. No. 0.0011;
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                                                                                                                                                                        Mismatches
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  Disulfide-bond
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Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  albumin fusion protein; cytostatic; anorectic; immunosuppressive; antidiabetic; antirheumatic; antiarthritic; psoriatic; cancer; non-Hodgkin's lymphoma; obesity; transplant rejection; psoriasis; type I diabetes mellitus; rheumatoid arthritis.
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Disulfide-bond
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Disulfide-bond
                                                                               N-PSDB; ABN87288.
                                                                                                              Rosen CA,
                                                                                                                                                                                                                25-OCT-2001.
                                                                                                                                                                                                                                    WO200179442-A2
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                                                                                                                                 HUMAN
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The present invention describes an albumin fusion protein (I) comprising a therapeutic protein: X and (a fragment or variant of) albumin
                                                                                                       New albumin fusion proteins, useful for treating diseases and disorders such as cancer, comprise therapeutic protein fused to albumin -
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25-APR-2000; 2000US-199384P.
21-DEC-2000; 2000US-256931P.
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                                                               Claim 1; Fig 11; 413pp; English
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278..289
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cc comprising a the fully defined sequence in ABB79006 of 585 amino acids, (where the fragment or variant has albumin or therapeutic protein: X c activity). (I) can have cytostatic, anorectic, immunosuppressive, antidiabetic, antirheumatic, antiarthritic and psoriatic activities. Cc antidiabetic, antirheumatic, antiarthritic and psoriatic activities. Albumin fusion proteins are stabilised therapeutic proteins e.g. cc and disorders such as non-Hodgkin's lymphoma, cancer, obesity, can disorders such as non-Hodgkin's lymphoma, cancer, obesity, cc transplant rejection, type I diabetes mellitus, rheumatoid arthritis c and psoriasis. Fusing albumin to therapeutic proteins stabilises the therapeutic proteins extends the in vitro or cin vivo biological activity. It also reduces the need to formulate c protein solutions with large excesses of carrier proteins to prevent container. The fusion proteins are easily dispensed with a simple c container. The fusion proteins are easily dispensed with a simple c formulation requiring minimal post storage manipulation. The fusion of therapeutic proteins to albumin confers stability in aqueous or other solution. The present sequence represents the mature human albumin (HA) crotein which is used in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; albumin; HA; fusion protein; immune system disorder; syphilis; transplant rejection; blood related disorder; myocardial infarction; hyperproliferative disorder; acute myeloid leukaemia; renal disorder; glomerulonephritis; cardiovascular disorder; arrhythmia; rhinitis; respiratory disorder; neurological disease; Alzheimer's disease; endocrine disorder; neurological disease; Alzheimer's disease; measles; gastrointestinal disorder; irritable bowel syndrome; HIV; human immunodeficiency virus; wound healing; renal cell carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human albumin (HA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                     melanoma; gene therapy; mutant; mutein.
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                                                                                                                                            Domain
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                                  Domain
                                                                    Domain
                                                                                                        Domain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                 170..176
                                                                                                                                                           /note= "Xaa represents the mutation of the
amino acid to any other amino acid"
                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                               /label= Loop_VII
                                                                                                                                                                                                                                                                                 /6..89
/label= Loop_II
                                                                                                                                                                                                                                               'label=_Loop_III
                                                                                                                         Label=
                                                                                                                                                                                                               label= Loop_IV
                                                                                                                                                                                                                                                                                                                         label= Loop_I
                 .abel=
                                                                               .abel= Loop_VI
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100.0%; Pred. No. 0.0
tive 0; Mismatches
                                                                                                            277
                                                                                                                   A_door
               LOOP_VIII
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AAE13399
                                                                                                                                                                  RESULT 48
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                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Fig 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Albumin fusion proteins comprising a therapeutic protein and albumin, useful in the treating metastatic renal cell carcinoma, metastatic melanoma, malignant melanoma, renal cell carcinoma, HIV (human immunodeficiency virus) or infection -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rosen CA,
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25-APR-2000; 2000US-199384P.
21-DEC-2000; 2000US-256931P.
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                                                                                                                                      AAE13399 standard; Protein; 585
                                                                                                                                                                                                                                                                                                                         Sequence
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Pred. No. (
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                                                                                                                                                                                                                                                                               0.0011;
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                                                                                                                                                                                                                                                                                             Length 585;
                                                                                                                                                                                                                                                                    Indels
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Human; albumin; HA; fusion protein; immune system disorder; transplant rejection; blood related disorder; myocardial ini hyperproliferative disorder; acute myeloid leukaemia; renal

infarction; nal disorder; syphilis; Human albumin (HA) protein

12-FEB-2002

(first entry)

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        The invention relates to albumin fusion proteins comprising therapeutic protein and human albumin (HA). The albumin fusion proteins are useful in the treatment, prevention, diagnosis, and/or detection of diseases, disorders such as immune system disorders (transplant rejection); blood related disorders (myocardial infarction), hyperproliferative disorders (childhood acute myeloid leukaemia); renal disorder (glomerulonephritis); cardiovascular disorders (arrhythnias); respiratory disorders (non-allergic rhinitis); neurological diseases (Alzheimer's disease); endocrine disorders (pheocytochroma); respiratory esystem disorders (syphilis); infectious diseases (measles); gastrointestinal disorders (irritable bowel syndrome) and wound healing. The albumin fusion proteins are also used in the treatment of metastated renal cell
                                                                                                                                                                                                                       Albumin fusion proteins comprising a therapeutic protein and albumin, useful in the treating metastatic renal cell carcinoma, metastatic melanoma, malignant melanoma, renal cell carcinoma, HIV (human immunodeficiency virus) or infection -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       respiratory disorder; neurological disease; Alzheimer's disease; endocrine disorder; pheocytochroma; reproductive system disorder; measles; gastrointestinal disorder; irritable bowel systemome; HUY; human immunodeficiency virus; wound healing; renal cell carcinoma;
                                                                                                                                                                                                                                                                                                                                      Rosen
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25-APR-2000; 2000US-199384P.
21-DEC-2000; 2000US-256931P.
proteins are carcinoma, me
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                                                                                                                                                                                             1; Fig 9; 325pp; English.
                                                                                                                                                                                                                                                                                                                                                                  PRINCIPIA PHARM CORP
                                                                                                                                                                                                                                                                                                                                                                                 HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene therapy.
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560..566
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76..89
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'label
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             label=
 melanoma,
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                                                                                                                                                                                                                                                                                                                                      Turner AJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                arrhythmia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                rhinitis;
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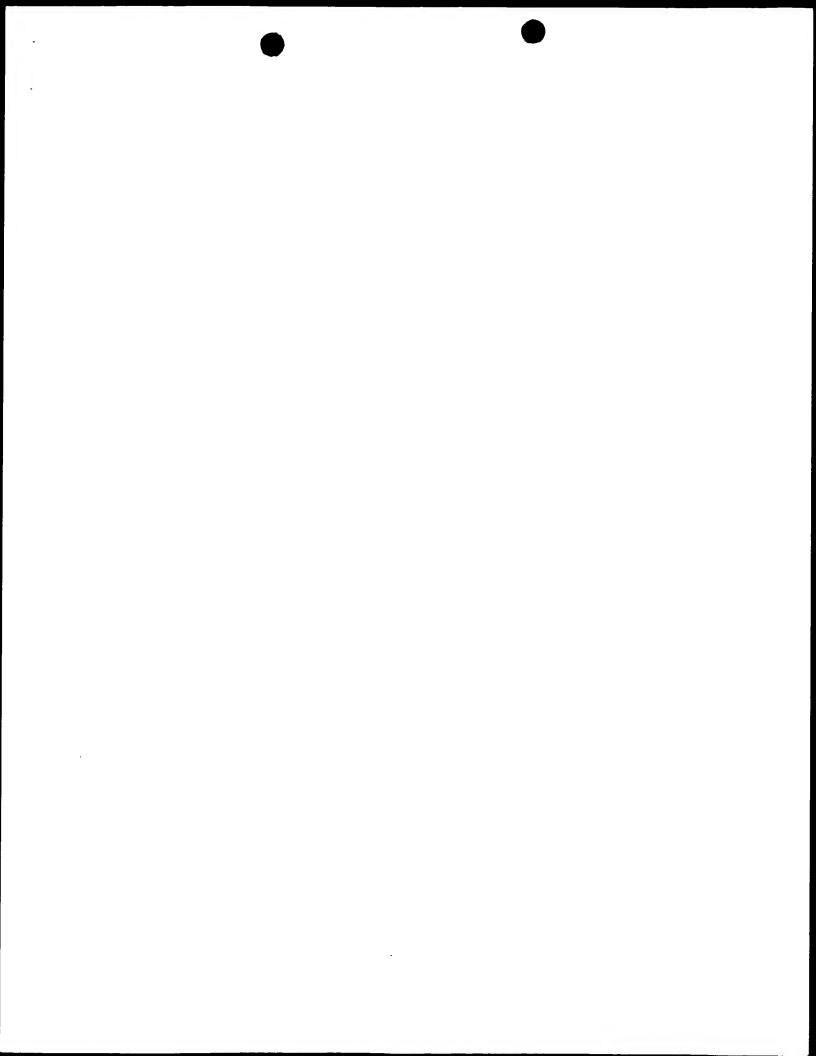
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RESULT 49
AAM52567
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                                         rejection), blood related disorders (e.g. myocardial infarction), hyperproliferative disorders (e.g. childhood acute myeloid leukemia), renal disorders (e.g. glomerulonephritis), cardiovascular disorders (e.g. arrhythmias), respiratory disorders (e.g. non-allergic rhinitis), neurological diseases (e.g. Alzheimer's disease), endocrine disorders (e.g. phecoytochroma), reproductive system disorders (e.g. syphilis), reproductive system disorders (e.g. syphilis),
                                                                                                                                         a therapeutic protein and albumin. The present sequence is the protein sequence for mature human serum albumin (HA), which was used to generate the fusion proteins of the present invention. The albumin fusion proteins are useful in the treatment, prevention, diagnosis, and/or detection of diseases/disorders such as immune system disorders (e.g. transplant diseases/disorders such as immune system disorders (e.g. transplant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; serum albumin; HA; antiinflammatory; immunosuppressive; cardiant; nootropic; neuroprotective; gene therapy; immune disorder; wound healing; hyperproliferative disorder; renal disorder; cardiovascular disorder; respiratory disorder; neurological disease; endocrine disorder;
                                                                                                                                                                                                                                                                                                            Albumin fusion proteins comprising a therapeutic protein and albumin, useful in the treating immune system disorders (e.g. transplant rejection), blood related disorders (e.g. myocardial infarction) and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunodeficiency virus) infection. Nucleic acid encoding albumin fusion
                                                                                                                                                                                                                                                                                            rejection), blood related dishyperproliferative disorders
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein is useful in gene therapy. The present sequence
                                                                                                                                                                                                                            The present invention relates to albumin fusion proteins,
                                                                                                                                                                                                                                                               Claim 1; Fig 15; 606pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                 Rosen CA,
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25-APR-2000;
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                               infectious diseases (e.g.
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; 2000US-199384P.
; 2000US-256931P.
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Pred. No.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                12-APR-2000; 2000US-229358P.
25-APR-2000; 2000US-199384P.
21-DEC-2000; 2000US-256931P.
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           (HUMA-) HUMAN GENOME SCI INC
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                                                                                                                                                                                                                              439..447
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                                                                                                                                                                                                                                                                                                                                                                           /label= Loop_II
                                                                                                                                                                   /label= Loop_XI
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                                                                                                                                            'label= Loop_XII
                                                                                                                                                                                                                                                                                                           label
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                                                                                                                                                                                                              label= Loop_IX
                                                                                                                                                                                                                                     label= Loop_VIII
                                                                                                                                                                                                                                                          label= Loop_VII
                                                                                                                                                                                       label= Loop_X
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                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Albumin fusion proteins comprising a therapeutic protein and albumin, useful in the treating immune system disorders (e.g. transplant rejection), blood related disorders (e.g. myocardial infarction) and
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                                                                                                                                                                                                                               Sequence
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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfIles1.pep:*
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     Gapext 0.5
                          US-08-153-799-14
US-08-448-196A-3
US-08-984-176-1
US-08-762-772-2
US-08-762-746-2
US-08-8-769-746-2
US-08-897-956A-2
PCTT-US-93-8-2
PCTS-08-93-9-68-2
US-08-97-689-2
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US-08-134-638-1
US-08-134-638-1
US-08-148-196A-4
US-08-186-409-13
US-08-448-196A-6
US-08-488-196A-6
US-08-488-196A-6
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US-08-883-364-1
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Sequence 17, Appli Sequence 6, Appli Sequence 6, Appli Sequence 3, Appli Sequence 34, Appli Sequence 9, Appli Sequence 9, Appli	equence 1, equence 1, equence 15, equence 17, equence 17, equence 17, equence 17, equence 17, equence 15, equence	quence 2, quence 10, quence 13, quence 2, quence 5, quence 4, quence 15, quence 11, quence 11	equence 1 equence 1 equence 1 equence 3 equence 3 equence 3 equence 1 equence 1	Sequence 10, Appli Sequence 1, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 28, Appli Sequence 15, Appl Sequence 11, Appli Sequence 11, Appli

ALIGNMENTS

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US-08-153-799-14

(US-08-153-799-14

; Sequence 14, Application US/08153799

; Patent NO. 5766883

; Patent INPORMATION:
   APPLICANT: Ballance, David J
   CORRESPONDENCE ADDRESS:
   NUMBER OF SEQUENCES: 23
   CORRESPONDENCE ADDRESS:
   ADDRESSEE: R Hain Swope, BoC Health Care Independent Country: USA STREET: 100 Mountain Avenue
   CITY: Murray Hill
   STATE: New Jersey
   COUNTRY: USA
   ZIP: 07974
   COMPUTER READABLE FORM:
   MEDIUM TYPE: Floppy disk
   COMPUTER: IBM PC Compatible
   OPERATING SYSTEM: PC-DOS/MS-DOS
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US-08-448-196A-3
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                              tches
COMPUTER READABLE FORM
                                                                                                                                                      APPLICANT: CARTER, DANIEL C.
TITLE OF INVENTION: BIOLOGIC
TITLE OF INVENTION: CONTAINI
TITLE OF INVENTION: RELATED
                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORIGINAL SOURCE:
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/847975
FILING DATE: 06-MAR-1992
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: 11
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NAME: Swope, R Hain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 29-APR-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/GB90/00650
FILING DATE: 26-APR-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: Region LOCATION: 1..585 OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: Region
LOCATION: 369..419
OTHER INFORMATION: ,
OTHER INFORMATION: ,
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                                                     STATE:
                                                                    STREET: MARSHALL SPACE FLIGHT CENTER CITY: HUNTSVILLE
                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (908) 771 6159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Swope, R Hain
REGISTRATION NUMBER: 24864
REFERENCE/DOCKET NUMBER: 9
                                                                                                       ADDRESSEE: NASA
                                                                                                                                                                                                                                                                                                                                                        1 DAHKSEVAHRFK 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 07/775952 FILING DATE: 29-OCT-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: GB 8909916.2
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                  35812
                                                  ALABAMA
                                 USA
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                                                                                                                                                       BIOLOGICALLY ACTIVE PROTEIN FRAGMENTS CONTAINING SPECIFIC BINDING REGIONS OF SERUM ALBUMIN OR RELATED PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 64; DB 1; 100.0%; Pred. No. 0.00049;
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natural HSA"
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Sequence 2, Application US/08702572

Ratent No. 5965386
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Kerry-Williams, Sean M
APPLICANT: Gilbert, Sarah C
TITLE OF INVENTION: Yeast Strains and Modified Albumins
                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: OXYGEN-TRANSPORTING ALBUMIN-BASED BLOOD REPLACEMENT TITLE OF INVENTION: COMPOSITION AND BLOOD VOLUME EXPANDER FILE REFERENCE: 08/984,176
CURRENT APPLICATION NUMBER: US/08/984,176
CURRENT FILING DATE: 1997-12-03
NUMBER OF SEQ ID NOS: 1
SOFTWARE: Patentin Ver. 2.0
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: CARTER, DANIEL C
APPLICANT: HO, JOSEPH X
APPLICANT: RUKER, FLORIAN
                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                      Query Match
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INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 585
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: BROAD JR., ROBERT L.
REGISTRATION NUMBER: 18,757
REFERENCE/DOCKET NUMBER: XX.
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/448,196A
FILING DATE: 23-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
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Local Similarity 100.0%;
es 12; Conservation
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100.0%; Pred. No. 0.00049;
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Pred. No.
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CORRESPONDENCE ADDRESS:

ADDRESSEE:

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Centeon L.L.C. 20 First Avenue NUMBER OF SEQUENCES:

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                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2, Application US/08769746 Patent No. 6274305 GENERAL INFORMATION:
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APPLICATION NUMBER: WO 95/23857
FILING DATE: 1.MAR-1995
APPLICATION NUMBER: GB 9404270.2
FILING DATE: 5-MAR-1994
ATTORNEY/AGENT INFORMATION:
NAME: NAOM! BISWAS
RECISTRATION NUMBER: 38,384
REFERENCE/DOCKET NUMBER: CE0114 US
TELEPHONE: 610/878/4294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: MicroSoft Word 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/702,5
FILING DATE: 11-NOV-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 2:
                               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMMBER: US/08/769,746
FILING DATE: 19-DEC-1996
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 585 amino acid
                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Sonnenschein, Carlos
APPLICANT: Soto, Ana M.
TITLE OF INVENTION: Inhibiting Proliferation of Cancer Cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                        STREET: 220 Montgome
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX:
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                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
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Carroll, Peter G
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                                                                                                                                                                                                                                                                                             220 Montgomery Street, Suite 2200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       585 amino acids
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                                                                                                                                                                                                                                     United States of America
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US-08-222-619-3
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                                                                                     Query Match
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INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                           CLASSIFICATION: 435
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
CENTRE: 609 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/222,619 FILING DATE:
                                                                                                                                                                                                                                                                                                                                           COUNTRY 2120-1789
ZIP: 91320-1789
COMPUTER READABLE FORM:
MEDITIM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: MB TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 705-8410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                  TOPOLOGY: un MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Wright, Samuel
TITLE OF INVENTION: Afamin: A Human Serum Albumin-Like
TITLE OF INVENTION: Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
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25
                                                    Y Match 100.0%;
Local Similarity 100.0%;
108 12; Conservative 0
                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 1840 DeHavilland Drive CITY: Thousand Oaks
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER:
                                                                                                                                                             STRANDEDNESS: unknown
                                                                                                                                                                                  TYPE:
                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U.S.
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               1 DAHKSEVAHRFK 12
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DAHKSEVAHRFK 36
                                                                                                                                                                               amino acid
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                                                                                                                                                                                             609 amino acids
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                                                                                                                                                    unknown
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                                                                                                                                  protein
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100.0%; Pred. No.
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                                                                          Score 64; DB 1;
Pred. No. 0.00051
                                                           Mismatches
                                                                          0.00051;
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                                                                                        Length 609;
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RESULT 7
US-08-433-037-4
; Sequence 4, Application US/08433037
; Patent No. 5707828

GENERAL INFORMATION:

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                                                                                                 SOFTWARE:
SEQ ID NO 2
LENGTH: 60
TYPE: PRT
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TELEPAX: (516) 742-4366
TELEX: 230 901 SANS UR
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 609 amino acids
                                                                                                                                                                                                                                                                                                                                                                          Sequence 2, Application US/08897956A Patent No. 6423512 GENERAL INFORMATION:
Best Local Similarity
                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                      FILE REFERENCE: 600-7244/CPA
CURRENT APPLICATION NUMBER: US/08/897,956A
CURRENT FILING DATE: 1997-07-21
PRIOR APPLICATION NUMBER: 60/022,689
PRIOR FILING DATE: 1996-07-26
NUMBER OF SEQ ID NOS: 38
NUMBER OF SEQ ID NOS: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                             08-897-956A-2
                                                                                                                                                                                                                                                                                                                    APPLICANT: Mary Ellen Digan
APPLICANT: Philip Lake
APPLICANT: Hermann Gram
                                                                                                                                                                                                                                                                                               APPLICANT: Hermann Gram
TITLE OF INVENTION: Fusion Polypeptides
                                                                             ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Diciglio, Frank S.
REGISTRATION NUMBER: 31,346
REFERENCE/DOCKET NUMBER: 9108Z
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: Floppy disk
COMPUTER: PATENTIAL PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US/08/433,037
FILING DATE: 03-WAY-1995
CLASSIFICATION: 435
CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Tschopp, Juerg F.
TITLE OF INVENTION: EXPRESSION OF HUMAN SERUM ALBUMIN IN
TITLE OF INVENTION: PICHIA PASTORIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 400 Garden City Plaza
CITY: Garden City
STATE: New Variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25 DAHKSEVAHRFK 36
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Local Similarity 100.0%;
les 12; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 DAHKSEVAHRFK 12
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                                                                                                                      609
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                                                                                                                                                              FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Brierley, Russell A.
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100.0%;
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Score 64; DB 4;
Pred. No. 0.00051;
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Pred. No.
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                Length 609;
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US-08-797-689-2
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 609 amino acids
                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                          APPLICANT: Fleer, Reinhard
APPLICANT: Fournier, Alain
APPLICANT: Guitton, Jean-Dominique
APPLICANT: Jung, Gerard
APPLICANT: Yeh, Patrice
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 91320-1789

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: FREE PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #11.0, Version #1.25

CURRENT APPLICATION DATA:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: AMGEN INC.
TITLE OF INVENTION: Afamin:
TITLE OF INVENTION: Protein
                                                                                                                                                                               TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: protein
                                                                               STREET: 500 Arcola
CITY: Collegeville
STATE: PA
                                                COUNTRY: US
ZIP: 19426
                                                                                                                                                                                                                                                                                                                                                                                                                                  25
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STREET: 1840 DeHavilland Drive
CITY: Thousand Oaks
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: PCT/US95/04075 FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 DAHKSEVAHRFK 12
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                                                                                                                 E: Rhone-Poulenc Rorer Inc
500 Arcola Road, 3C43
                                                                 USA
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                                                                                                                                                                               NOVEL BIOLOGICALLY ACTIVE POLYPEPTIDES, PREPARATION THEREOF AND PHARMACEUTICAL COMPOSITION CONTAINING SAID POLYPEPTIDES
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Pred. No.
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OPERATING SYSTEM: SYSTEM SOFTWARE: WORD 5.1 (PACURRENT APPLICATION DATA:

System 7.1

31-JAN-1997

US/08/797,689 (PatentIn)

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US-08-256-938-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 28-JAN-1993
ATTORNEY/AGENT INFORMATION:
NAME: Smith ph.D., Julie K.
REGISTRATION NUMBER: p-38,619
REFERENCE/DOCKET NUMBER: ST92006-US
TELECOMMUNICATION INFORMATION:
TELEPAX: (610) 454-3839
TELEFAX: (610) 454-3809
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 610 amino acide
TYPE: amino acide
TYPE: amino acide
TYPE: amino acide
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APPLICATION UNMER: US/08/256,938
FILING DATE: US/08/256,938
FILING DATE: 435
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: FR 92/01065
APPLICATION NUMBER: FR 92/01065
APPLICATION NUMBER: 31-JAN-1992
ATTORNEY/AGENT INFORMATION:
NAME: GOOdman, Rosanne
REGISTRATION NUMBER: 32,534
REFERENCE/DOCKET NUMBER: ST92007-US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (610) 454-3817
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Yeh, P.
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APPLICATION NUMBER: US 08/256,927
FILING DATE: 28-JUL-1994
                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
OPERATING SYSTEM: System 7.1
SOFTWARE: Word 5.0 (Patentin)
CURRENT APPLICATION DATA:
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ADDRESSEE: Rhone-Poulenc Rorer Inc.
STREET: 500 Arcola Road, 3C43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Yeh, Patrice
TITLE OF INVENTION: NEW POLYPEPTIDES HAVING GRANULOCYTE
TITLE OF INVENTION: COLONY STIMULATING ACTIVITY, PREPARATION THEREOF AND
TITLE OF INVENTION: PHARMACEUTICAL COMPOSITIONS CONTAINING SAID POLYPEPTIDES
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/FR93/00085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 500 ALCULO CITY: Collegeville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25 DAHKSEVAHREK 36
                                                                                                                                                                                                                                                                                                                                                                                                          ZIP:
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                                                                                                                                                                                                                                                                                           (PatentIn)
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RESULT 13 US-08-797-689-16

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GENERAL INFORMATION:
APPLICANT: Yeh, P
                                                                  Best Loc
Matches
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 783 amino acids
                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: Macintosh

OPERATING SYSTEM: System 7.1

SOFTWARE: Word 5.0 (Patentin)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/256;
                                                                                                                                                                                                                                            TELEFAX: (610) 454-3808 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Yeh, Patr
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                          NAME: Goodman, Rosanne
REGISTRATION NUMBER: 32,534
REFERENCE/DOCKET NUMBER: ST
TELECOMMUNICATION INFORMATION:
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ADDRESSEE: Rhone-Poulenc Rorer Inc
STREET: 500 Arcola Road, 3C43
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                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 787 amino acids
                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 31-JAN-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                         MOLECULE TYPE: protein
203
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TOPOLOGY: linear
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STATE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 19426
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                                                                                 Local Similarity
                                                                                                                                                                         TOPOLOGY:
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                               1 DAHKSEVAHRFK 12
DAHKSEVAHREK 214
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XGY: linear
                                                                100.0%; Score 64; DB 1; ilarity 100.0%; Pred. No. 0.00067; Conservative 0; Mismatches 0;
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100.0%; Pred. No. 0.00067;
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                                                                                                                                                   GENERAL INFORMATION:
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APPLICATION NUMBER: US 08/256,927

FILING DATE: 28-JUL-1994

APPLICATION NUMBER: FR 92/01064

FILING DATE: 31-JAN-1992

PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/FR93/00085

FILING DATE: 28-JAN-1993

ATTORNEY/AGENT INFORMATION:

NAME: Smith Ph.D., Julie K.

REGISTRATION NUMBER: P-38,619

REFERENCE/DOCKET NUMBER: ST92006-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 16,
Patent No. 5
             APPLICANT: Mary Ellen Digan
APPLICANT: Philip Lake
APPLICANT: Hermann Gram
TITLE OF INVENTION: Fusion Polypeptides
FILE REFERENCE: 600-7244/Cpa
CURRENT APPLICATION NUMBER: US/08/897,956A
CURRENT FILING DATE: 1997-07-21
PRIOR APPLICATION NUMBER: 60/022,689
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 787 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
8-797-689-16
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APPLICANT:
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CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: NOVEL BIOLOGICALLY ACTIVE POLYPEPTIDES,
TITLE OF INVENTION: PREPARATION THEREOF AND PHARMACEUTICAL COMPOSITION
TITLE OF INVENTION: CONTAINING SAID POLYPEPTIDES
NUMBER OF SEQUENCES: 36
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
TOPOLOGY: linear
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OPERATING SYSTEM: System 7.1
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5876969
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500 Arcola Road, 3C43
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Yeh, Patrice
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Guitton, Jean-Dominique
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US-08-134-638-1
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                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 1996-07-26
NUMBER OF SEQ ID NOS: 38
SOFTWARE: FastSEQ for Windows
SEQ ID NO 3
                                                                          Best
                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                TELEFAX: (602) 947-2663 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Strand, F
TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE: OTHER INFORMATION: Fusion polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                            NAME: Weiss Harry M
REGISTON NUMBER: 19,497
REFERENCE/DOCKET NUMBER: 17:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (602) 994-8888
                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 10/12/93
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRI: 5.221
ZIP: 85082-4321
COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
TYPE: Diskette, 5.25 inch, 1.2 Mb storage
                                                                                                                                                                                                                                                                                                                         FILING DATE: N/A ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                212 DAHKSEVAHREK 223
                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: sin
                                                                       Local
                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: N/A
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nes 12; Conserv
                                                                                                                                                                                 LENGTH:
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DTHKSEIAHRFK 12
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                                                         l Similarity
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                                                          Conservative
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                                                       Score 59; DB 1; Length 582; Pred. No. 0.004; 1; Mismatches 1; Indels
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Pred. No. 0.00085;
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US-08-448-196A-4
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US-08-448-196A-7
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                                                                                                                                                                                                                                             Sequence 7, Application US/08448196A
Patent No. 5780594
GENERAL INFORMATION:
APPLICANT: CARTER, DANIEL C.
TITLE OF INVENTION: BIOLOGICALLY ACTIVE PROTEIN FRAGMENTS
TITLE OF INVENTION: CONTAINING SPECIFIC BINDING REGIONS OF SERUM ALBUMIN OR
TITLE OF INVENTION: RELATED PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local Similarity 83.3%;
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INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 583 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: CARTER, DATTILE OF INVENTION: ETITLE OF INVENTION: FITTLE OF INVENTION: FUNDMBER OF SEQUENCES:
                                                                                                                                                                                                     TITLE OF INVENTION: BIOI TITLE OF INVENTION: CONTITLE OF INVENTION: RELAUNDBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 18,757
REFERENCE/DOCKET NUMBER: XX.
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Ve
CUREBY APPLICATION DATA:
APPLICATION UNDBER: US/08/448,196A
FILING DATE: 23-MAY-1995
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MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: BROAD JR., ROBERT L.
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 583 amino acids
TYPE: amino acid
TOPOLOGY: linear
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                                                                                                                       STREET: MARSHALL SPACE FLIGHT CENTER CITY: HUNTSVILLE STATE: ALABAMA
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                                                                                 ZIP: 35812
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                                                                                                      COUNTRY:
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                                                                                                        USA
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BIOLOGICALLY ACTIVE PROTEIN FRAGMENTS

CONTAINING SPECIFIC BINDING REGIONS OF SERUM ALBUMIN OR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RELATED PROTEINS
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NAME: BROAD JR., ROBERT L.
REGISTRATION NUMBER: 18,757
REFERENCE/DOCKET NUMBER: XX/M
TELECOMMUNICATION INFORMATION:
TELEPHONE: 205-544-0021
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APPLICANT: FASANO,
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INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 584 amino acids
                                                                                                   CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
ANAME: KIT, Gordon
REGISTRATION NUMBER: 30,764
REFERENCE/DOCKET NUMBER: A-6888
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 293-7860
TELEPHONE: (202) 293-7860
                                                INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/0
FILING DATE: 23-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 20 FEB
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   OLECULE TYPE:
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Local Similarity 83.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Washington, D.C.
                TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                         FILING DATE
                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP:
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peptide
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Pred. No. 0.004;
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                                               Patent No. 5948629
GENERAL INFORMATION:
                                                                      Sequence 13, Application US/09186409 Patent No. 5948629
                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (202) 293-7060
TELEPAX: (202) 293-7860
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acids
NUMBER OF SEQUENCES:
         APPLICANT: FASANO, Alessio
TITLE OF INVENTION: ZONULA OCCLUDENS TOXIN RECEPTOR
                                                                                                                                                                                                                                                                                                                                             HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: A-TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC-compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
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NAME: KIT, Gordon
REGISTRATION NUMBER: 30,764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSEE: SUGHRUE, MION, ZINN, MACPEAK & SEAS
STREET: 2100 Pennsylvania Avenue, N.W., Suite 800
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APPLICATION NUMBER: (
FILING DATE: 20 FEB :
CLASSIFICATION:
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TITLE OF INVENTION: ZONULA OCCLUDENS TOXIN RECEPTOR
                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/01 FILING DATE: 17 FEB 1998
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                                                                                                                                                                                 1 AHKSEVAHRFK 11
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5912323
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                                                                                                                                                                                                                                                                   90.6%; Score 58; DB 2; L
100.0%; Pred. No. 9.5e-05;
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100.0%; Pred. No. 9.5e-05;
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                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                  APPLICANT: CARTER, I
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
HYPOTHETICAL:
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APPLICATION NUMBER:
FILING DATE: 17 FEB
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Re-Lease #1.0, Version #1.25
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TRM PC compatil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (202) 293-7060
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: SUGHRUE, MION, ZINN, MACPEAK & SEAS
                                                                                                                                                     STATE:
                                                                                                                                                                    STREET: MARSHALL SPACE FLIGHT CENTER CITY: HUNTSVILLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 2100 Penns
CITY: Washington,
 APPLICATION NUMBER:
                                                                                                                     ZIP:
                                                                                                                                  COUNTRY: USA
                                                                                                                                                                                                     ADDRESSEE: NASA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 100 les 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 30 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: FILING DATE: 5 NOV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                             1 AHKSEVAHRFK 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 AHKSEVAHREK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH:
                                                                                                                     35812
                                                                                                                                                    ALABAMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               D.C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
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NO
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                                                                                                                                                                                                                                                    DANIEL C.
BIOLOGICALLY ACTIVE PROTEIN FRAGMENTS
CONTAINING SPECIFIC BINDING REGIONS OF SERUM ALBUMIN OR
RELATED PROTEINS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90.6%; Score 58; DB 2;
100.0%; Pred. No. 9.5e-0
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US/08/448,196A
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CLASSIFICATION:

23-MAY-1995

Query Match

Local Similarity

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US-08-448-196A-6
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                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PALENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/448,196A
FILING DATE: 23-MAY-1995
CLASSIFICATION: S30
ATTORNEY/AGENT INFORMATION:
NAME: BROAD JR., ROBERT L.
REGISTRATION NUMBER: 18,757
REGISTRATION NUMBER: 18,757
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                                                                                                                             TELEPHONE: 205-544-0021
TELEPAX: 205-544-0258
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 583 amino acide
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TITLE OF INVENTION: BIOLOGICAL
TITLE OF INVENTION: CONTAINING
TITLE OF INVENTION: RELATED PI
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
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INFORMATION FOR SEQ ID NO: 5:
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                                  MOLECULE TYPE: p:
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                        REGISTRATION NUMBER: 18,757
REFERENCE/DOCKET NUMBER: XX/
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: protein HYPOTHETICAL: NO ANTI-SENSE: NO FRAGMENT TYPE: N-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 583 amino acid
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REFERENCE/DOCKET NUMBER: XX/MFS-28402-2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT
                  FRAGMENT TYPE:
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STREET: MA
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TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE:
                                                                                           TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 DTHKSEIAHRF 11
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BROAD JR., ROBERT L.
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                    N-terminal
                                                                        protein
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CONTAINING SPECIFIC BINDING REGIONS OF SERUM ALBUMIN OR
RELATED PROTEINS
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    Mismatches

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Pred. No. 0.033;
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Sequence 12, Application US/09024198
; Batent No. 5912323
; GENERAL INFORMATION:
APPLICANT: FASANO, Alessio
TITLE OF INVENTION: ZONULA OCCLUDENS TOXIN RECEPTOR
NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSE: SUGHRUE, MION, ZINN, MACPEAK & SEAS
ADDRESSEE: SUGHRUE, MION, ZINN, MACPEAK & SEAS
STREET: 2100 Pennsylvania Avenue, N.W., Suite 800
                                                                                                                                                                                    RESULT 24
US-09-024-198-12
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (202) 293-7860 INFORMATION FOR SEQ ID NO: (SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
ADDRESSES: SUGHRUE, MION, ZINN, MACPEAK & SEAS
STREET: 2100 Pennsylvania Avenue, N.W., Suite 800
CITY: Washington, D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 30,764
REFERENCE/DOCKET NUMBER: A-
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: peptide HYPOTHETICAL: NO
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NAME: KIT, Gordon
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                          y Match 78.1%;
Local Similarity 90.0%;
les 9; Conservative
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FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
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CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                               Score 50; DB 2; Length 13; Pred. No. 0.0028;
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Patent No. 5948629
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 12
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/186,409
FILING DATE: 5 NOV 1998
                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: FASANO, Alessio
TITLE OF INVENTION: ZONULA OCCLUDENS TOXIN RECEPTIVE OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSEE: SUGHRUE, MION, ZINN, MACPEAK & SEAS
ADDRESSEE: SUGHRUE, MION, ZINN, MACPEAK & SEAS
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/8
FILING DATE: 20 FEB 1997
                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/0
FILING DATE: 17 FEB 1998
CLASSIFICATION:
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REFERENCE/DOCKET NUMBER: A-6988
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 293-7060
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APPLICATION NUMBER: 08/8
FILING DATE: 20 FEB 1997
CLASSIFICATION:
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CITY: Washington,
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mes 9; Conserv
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                                                                                                                                  CLASSIFICATION:
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90.0%;
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                 08/803,364
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                                                                                            09/024,198
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US-08-803-364-3
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                                                                 US-08-803-364-1
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Patent No. 5864014
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 Best Local Similarity Matches 9; Conserv
                               Query Match
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                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: KIT, Gordon
REGISTRATION NUMBER: 30,764
REFERENCE/DOCKET NUMBER: A-68
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 293-7060
                                                                            TYPE: amino a
TOPOLOGY: lir
MOLECULE TYPE:
HYPOTHETICAL: 1
                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
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MOLECULE TYPE:
HYPOTHETICAL:
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REGISTRATION NUMBER: 30,764
REFERENCE/DOCKET NUMBER: A-T
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 293-7060
                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 16 amino acids
                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/
FILING DATE: 20 FEB 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 2100 CCITY: Washington, D.C.
STATE: D.C.
STATE: T.C.A.
                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/803,364 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: SUGHRUE, MION, ZINNY, FINCE STREET: 2100 Pennsylvania Avenue, N.W., Suite 800
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les 9; Conserv
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                                                                                                                            amino acid
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   Conservative
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NO
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NO
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ZONULA OCCLUDENS TOXIN RECEPTOR
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90.0%;
                 78.1%;
90.0%;
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Score 50; DB 2; Length 16; Pred. No. 0.0035; 0; Mismatches 1; Indels
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Pred. No. 0.0028;
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                                                                                                                                                                                                                                      RESULT 28
                                                                                                     Sequence 10, Application US/09186409
PATENT NO. 5948629
GENERAL INFORMATION:
APPLICANT: FASANO, Alessio
TITLE OF INVENTION: ZONULA OCCLUI
NUMBER OF SEQUENCES: 18
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (202) 293-706
TELEPAX: (202) 293-7860
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: ZONULA OCCLUDENS TOXIN KELEFIUM NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSE: SUGHRUE, MION, ZINN, MACPEAK & SEAS
STREET: 2100 Pennsylvania Avenue, N.W., Suite 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
CRASSIFICATION DATA:
PRIOR APPLICATION DATA:
08/803,364
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/024,198
FILING DATE: 17 FEB 1998
             CORRESPONDENCE ADDRESS:
ADDRESSEE: SUGHRUE, MION, ZINN, MACPEAK & SEAS
STREET: 2100 Pennsylvania Avenue, N.W., Suite 800
CITY: Washington, D.C.
STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: A-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 293-7060
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: KIT, Gordon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: ami
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE: 20 FEI CLASSIFICATION:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 2100 Pennsylva
CITY: Washington, D.C.
STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
nes 9; Conserv
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ZIP: 20037
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GY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16 amino acids
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U.S.A.
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                                                                                                     ZONULA OCCLUDENS TOXIN RECEPTOR
                                                                                                                                                                                                                                                                                                                                                               78.1%; Score 50; DB 2; Length 16; 90.0%; Pred. No. 0.0035; tive 0; Mismatches 1; Indels
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Patent No. 5550114
GENERAL INFORMATION:
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Best Local Similarity 90.0%;
Matches 9; Conservative
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INFORMATION FOR SEQ ID NO:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
ATTORNEY/AGENT INFORMATION:
NAME: Zivin, No. 5550114man H.
REGISTRATION NUMBER: 25,385
REFERENCE/DOCKET NUMBER: 2560/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 977-9550
                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.24
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 16 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 30,764
REFERENCE/DOCKET NUMBER: A.-
TELECOMMUNICATION INFORMATION.
TELEPHONE: (202) 293-7060
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APPLICATION NUMBER:
FILING DATE: 20 FEB
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                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: COOPER & DUNHAM
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Strayer, TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
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                                                                                                               APPLICATION NUMBER: UPFILING DATE: 19930402
CLASSIFICATION: 514
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FILING DATE: 5 NOV 1998
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                            CITY: New York STATE: New Yorl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 1
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                        COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                            EPIDERMAL GROWTH FACTOR INHIBITOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                David S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09/024,198
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                                         2560/42559
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TELEFAX:

(212) 977-9809

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RESULT 31
US-08-469-856-9
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                                                                                                                                                                                                                                                                      Sequence 9, Application US/08469856 Patent No. 5650307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 7, Application US/08530340A Patent No. 6255452
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/08/530,340A CURRENT FILING DATE: 1995-12-22 NUMBER OF SEQ ID NOS: 7
                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Strayer, David S
TITLE OF INVENTION: Epidermal Growth Factor Inhibitor
FILE REFERENCE: JEFF-0226
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: An epidermal OTHER INFORMATION: growth factor inhibitor peptide
                                                                                                                                                                                 APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                             APPLICANT: Schrammeijer, Barbara
APPLICANT: Verwoerd, Tewnis C.
APPLICANT: VandenElzen, Peturs J.M.
TITLE OF INVENTION: PRODUCTION OF HETEROLOGOUS PROTEINS
TITLE OF INVENTION: PLANTS AND PLANT CELLS
                                                                             NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENGTH: 10
                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
COUNTRY:
                                            STREET:
                          CITY: Washington
                                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                             1 DTHKSEIAHR 10
                                                                                                                                                                                                                                                                                                                                                                                                            1 DAHKSEVAHR 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 DAHKSEVAHR 10
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                                              2000 Pennsylvania Avenue, NW, Ste. 5500
                                                                                                                                                                                               Hoekema, Andreas
Dekker, Bernardus M.M.
                                                                                                                                                                                                                                  Sijmons,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                               MORRISON & FOERSTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                            75.0%;
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                                                                                                                                                                                                                                      Peter C.
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Pred. No. 0.0048;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 48; DB 1;
Pred. No. 0.0048;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 10;
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Best Local !
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/469,856
FILING DATE: 06-UN-1995
CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
NUMBE: MITBRASTIGE KATE H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
INFORMATION FOR SEQ ID NO:
                                                                                                                                    FILING DATE: 19910708
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE P 4021571.7
FILING DATE: 06-JUL-1990
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: BURTSCHER, Helmut
APPLICANT: MOLLERING, Hans
TITLE OF INVENTION: CLONED N-METHYLHYDANTOINASE
                                               NAME: Murray, Robert B.
REGISTRATION NUMBER: 22,980
REFERENCE/DOCKET NUMBER: 91
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 659-2930
                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: MURASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFERENCE, DOCKET NUMBER: 61
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                TTTEPAX: (202) 887-0357
                                                                                                                                                                                                                             APPLICATION NUMBER: US
FILING DATE: 19910708
                                                                                                                                                                                                                                                                                                                                                                                          STREET: 1725 K Street, N.W., Suite 1000 CITY: Washington D.C. COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (202) 80, ----
TELEX: 90-4030 MRSNFOERSWSH
TO TO NO: 9:
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                                                                                                                                                                                                                                                                                                                                                                       ZIP: 20006
                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Armstro
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les 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            I: 11 amino acids
amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Armstrong, Nikaido, Marmelstein, Kubovcik &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (202)887-1500
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                                                                                      911014
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                                                                                                                                                                                                                                                                                    Version #1.25
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0;
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US-08-258-614-2
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                                                                                                                                     , MOLECULE TYPE: protein US-08-258-614-2
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                                                                                                    Query Match
                                                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US/08/021,856
APPLICATION NUMBER: US/08/021,856
APPLICATION DATA:
APPLICATION DATA:
APPLICATION UNMBER: DE P 40 21 571.7
FILING DATE: 06-7UL-1990
PRIOR APPLICATION DATA: US/07/727,814
APPLICATION UNMBER: US/07/727,814
APPLICATION UNMBER: US/07/727,814
APPLICATION TIPORMATION:
NAME: Kitts, Monica C.
REGISTRATION NUMBER: 9564-3007
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)638-5000
                                                                                                                                                                                                                                               TELEFAX: (202)638-4810
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            quence 2, Application US/08258614
ent No. 5432070
NUERAL INFORMATION:
APPLICANT: SCHUMACHER, Gunther
                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 1288 amino acid
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/258,614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: CLONED N-METHYLHYDANTOINASE NUMBER OF SEQUENCES: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           577 DAHFEEYGHRFK 588
577 DAHFEEYGHRFK 588
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                                                         y match
Local Similarity 66.7%;
hes 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A. ZIP: 20005-5701
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: D.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Washington
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                               1 DAHKSEVAHRFK 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E: Nikaido, Marmelstein, Murray & Oram 655 Fifteenth Street N.W. Suite 330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1288 amino acids
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MOLLERING, Hans
                                                                                                                                                                                                             1288 amino acids
                                                                                                                                                                                                                                                               (202) 638-4810
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66.7%;
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                                                                    0; Mismatches
                                                                                   Score 43; I
Pred. No. 8.
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Pred. No. 8.2;
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                                                                                   DB 1;
8.2;
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                                                                    4; Indels
                                                                                                     Length 1288;
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; ORGANISM: Homo sapiens US-09-165-926-2
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RESULT 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 2, Application US/09165926 Patent No. 6461875 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 2
                                                                                                                         Query Match
Best Local Similarity 58.:
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Ver. SEQ ID NO 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Bar-Or, David
APPLICANT: Lau, Edward
APPLICANT: Lau, Edward
APPLICANT: Lau, Edward
APPLICANT: Minkler, James V.
TITLE OF INVENTION: Test for Rapid Evaluation of Ischemic States and Kit
TILE REFERENCE: ISC35269-183796
CURRENT APPLICATION NUMBER: US/09/165,926
CURRENT FILING DATE: 1998-10-02
NUMBER OF SEQ ID NOS: 3
                                                                                                                                                                                                                                                                                                                                               APPLICANT: Kurzbauer, Robert
APPLICANT: Schaffner, Gotthold
TITLE OF INVENTION: Chicken Embryo Lethal Orphan
FILE REFERENCE: 0652.1800000
CURRENT APPLICATION NUMBER: US/09/171,461
CURRENT FILING DATE: 1999-01-12
EARLIER APPLICATION NUMBER: PCT/EP97/01944
EARLIER FILING DATE: 1997-04-18
NUMBER OF SEQ ID NOS: 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Baker, Adam
APPLICANT: Cotten, Matthew
APPLICANT: Chiocca, Susanna
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.0
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TYPE: PRT
                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: CELO VIRUS
                                                                                                                                                                                                                        OTHER INFORMATION: Position: 6501..9866/Product: E2b pol
                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                 LENGTH: 1121
                                                       132 DSHPGEVARRFR 143
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                                                                                        1 DAHKSEVAHRFK 12
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o. 6335016
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                                                                                                                                              62.5%;
58.3%;
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100.0%; Pred. No.
tive 0; Mismatc
                                                                                                                             Score 40; DB Pred. No. 25; 2; Mismatches
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                                                                                                                                                                  Length 1121;
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                                                                                                                                   Indels
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; Sequence 15, Application US/08738168B; Patent No. 613298B; GENERAL INFORMATION: APPLICANT: Sugino, Hiromu; APPLICANT: Nakamura, Takanori

US-08-738-168B-15

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                                                                              ; ORGANISM: Staphylococcus epidermidis US-09-134-001C-5307
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                                                                                                                                       SEQ ID NO 5307
LENGTH: 80
                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: LYNN DOUGETTE-Stamm et al
APPLICANT: LYNN DOUGETTE-Stamm et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity Matches 7; Conserv
    Matches
                  Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                        FILE REFERENCE: GTC-007
CURRENT APPLICATION UMMBER: US/09/134,001C
CURRENT FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR FILING DATE: 1997-11-08
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5674
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION UNMBER: US/08/738,168B
FILING DATE: 25-OCT-1996
PRIOR APPLICATION DATA: APPLICATION DATA: JP 280939/1995
FILING DATE: 27-OCT-1995
PRIOR APPLICATION DATA: JP 174909/1996
APPLICATION UNMBER: JP 174909/1996
FILING DATE: 04-JUL-1996
ATTORNEY/AGENT INFORMATION:
NAME: DESCRIPT DATA!
                                                                                                                                                                                                                                                                                                                                                                                                                    equence 5307, Ap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPAX: 617-523-6440
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 514 amino acids
                                                                                                                       TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Resnick, David S.
REGISTRATION NUMBER: 34,235
REFERENCE/DOCKET NUMBER: 342/46901
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-523-3400
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Shouji, Hiroki
TITLE OF INVENTION: NEURONAL CELL-SPECIFIC RECEPTOR PROTEIN
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE: DIKE, BRONSTEIN, ROBERTS & CUSHMAN, LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                       -134-001C-5307
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                Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
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Similarity 6; Conserv
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    Conservative
                  56.2%;
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70.0%;
<u>ب</u>
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                Score 36; DB
Pred. No. 7.5;
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Pred. No. 25;
Mismatches
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                                  DB 4; Length 80;
<u>ب</u>
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Indels
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Gaps
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                                                                                                                                            RESULT 39
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Best Local S
Matches 7
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MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOSTWARE: PASENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/357,533A
FILING DATE: 16-DSC-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: CRP-073FW
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                       Sequence 11, Application US/08459009
                                                                    Patent No. 5861479
GENERAL INFORMATION:
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                                 APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: protein FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 513 amino acids
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APPLICANT: OPPERMANN, HERMANN
APPLICANT: KUBERASAMPATH, THANGAVEL
APPLICANT: SMART, JOHN E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: Protein LOCATION: 1..513 OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                               312 EGHKPSIAHRDFK 324
                                                                                                                                                                                                                                                                    Local Similarity
nes 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (506)-435-0992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 45 SOUTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: sin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: PATENT ADMINISTRATOR, CREATIVE BIOMOLECULES ADDRESSEE: INC
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OPPERMANN, HERMANN
KUBERASAMPATH, THANGAVEL
SMART, JOHN E
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                                                       DONALD F
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                                                                                                                                                                                                                                                                                  53.9%;
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Pred. No. 1.1e+02;
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                                                                                                                                                                                                                                                                                                   Length 513;
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APPLICANT:

NOVEL MORPHOGEN CELL SURFACE RECEPTOR

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US-08-459-951-11
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FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION 1435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/357,533
FILING DATE: 16-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: KELLY, ROBIN D
REGISTRATION NUMBER: CRP-073FW
TELECOMMUNICATION INFORMATION:
TELEPHONE: (508)-435-9001
                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: JIN, DONALD F
APPLICANT: OPDERMANN, HERMANN
APPLICANT: KUBERASAMPATH, THANGAVEL
APPLICANT: SMART, JOHN E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (508)-435-0992
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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COUNTRY: USA
ZIP: 01748
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                 NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                     TITLE OF INVENTION: NOVEL MORPHOGEN CELL SURFACE RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                         312 EGHKPSIAHRDFK 324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 513 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOT OCT.
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OTHER INFORMATION:
OTHER INFORMATION:
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                                                                                       STREET: 45 SOUTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 DAHKSEVAHR-FK 12
                                                                                                                               ADDRESSEE:
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                                                                                                              E: INC
45 SOUTH STREET
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linear
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                                                                                                                                                PATENT ADMINISTRATOR, CREATIVE BIOMOLECULES
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Patent No. 5831050
         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTION PC-DOS/MS-DOS
SOFTWARE: PATENTION DATA:
APPLICATION NUMBER: US/08/357,533A
FILING DATE: 16-DEC-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: KELLY, ROBIN D
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: CRP-073FW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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APPLICATION NUMBER: US 08/357,533
APPLICATION NUMBER: US 08/357,533
APPLICATION INFORMATION:
ATTORNEY AGENT INFORMATION:
NAME: KELLY, ROBIN D
REGISTRATION NUMBER: 34,637
APPLICATION NUMBER: 34,637
                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: NOVEL MORPHOGEN CELL SURFACE RECEPTOR NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
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TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: JIN,
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ELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 53.8 nes 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION: 1..513
OTHER INFORMATION:
OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                           STATE:
                                                                                                                                                                                                                                                                                                          CITY: HOPKINTON
                                                                                                                                                                                                                                                                                                                              STREET:
                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
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(508)-435-0992
                                                                                                                                                                                                                                                                          USA
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KUBERASAMPATH, THAI
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RECEPTOR"
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Pred. No. 1.1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 3;
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OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION: FUS-08-357-533A-12
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INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 536 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-459-009-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/357,533
FILING DATE: 16-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: KELLY, ROBIN D
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: CRP-073FW
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 12,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                          TELEFAX: (508)-435-099
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 536 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tent No.
                                                TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/459,009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 01748
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: NOVEL MORPHOGEN CELL SURFACE RECEPTOR NUMBER OF SEQUENCES: 12 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: OPDERMANN, HERMANN
APPLICANT: OPDERMANN, HERMANN
APPLICANT: KUBERASAMPATH, THANGAVEL
APPLICANT: SMART, JOHN E
                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               335 ECHKPSIAHRDFK 347
LOCATION:
                   NAME/KEY:
                                                                                                                                                                                                TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: HOPKINTON
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nes 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: PATENT ADMINISTRATOR, CREATIVE BIOMOLECULES ADDRESSEE: INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: Protein LOCATION: 1..536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 DAHKSEVAHR-FK 12
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5861479
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                                                                                                                      536 amino acids
Protein
1..536
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                                                                                                                                                                                (508)-435-0992
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                                                                                                                                                                                                    (508) -435-9001
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53.8%; Pred. No. 1.1e+02;
tive 2; Mismatches 3; Indels 1
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                                                                                                                                                                                                                                         CRP-073FW
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Query Match
Best Local Similarity
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                                                                                            US-08-459-951-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No. 6093547
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/357,533
FILING DATE: 16-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: KELLY, ROBIN D
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: CRP-073FW
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
Matches 7; Conserv
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                                                                                                                                                                                                                                                                  TELEFAX: (508)-435-0992
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/459,951
                                                                                                                                                                        MOLECULE TYPE:
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 01748
COMPUTER READABLE FORM:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       335 EGHKPSIAHRDFK 347
                                                                                                           OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                TYPE: STRANDEDNESS: SI
                                                                                                                                      NAME/KEY: Protein LOCATION: 1..536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: PATENT ADMINISTRATOR, CREATIVE BIOMOLECULES, ADDRESSEE: INC
                                                                                                                                                                                                                                                                                                                  TELEPHONE:
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1 DAHKSEVAHR-FK 12
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45 SOUTH STREET
                                                                                                                                                                                                                                                    536 amino acids
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OPPERMANN, HERMANN
KUBERASAMPATH, THANGAVEL
                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                  (508) -435-9001
                                                                                                                                                                                       protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Floppy disk
                                                                                                                                                                                                                  single
                                              53.9%;
53.8%;
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                                                                                                           /note= "HUMAN ACTIVIN TYPE RECEPTOR"
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                                            Score 34.5; DB 3;
Pred. No. 1.1e+02;
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                             Mismatches
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                                                           Length 536;
                               Indels
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RESULT 44
US-08-586-039B-37
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Patent No. 614007
                                                                                                                                                                                             Sequence 39, Application US/08586039B Patent No. 6140073
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -08-586-039B-37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/586,03:
FILING DATE: 16-JAN-1996
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/124,259
FILING DATE: 20-SEP-1993
APPLICATION NUMBER: 07/676,436
FILING DATE: 28-MAR-1991
ATTORNEY/ACENT INFORMATION:
NAME: Hand, J. Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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lest Local Similarity 54.5%;
latches 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC COMPATIBLE
COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 6
CURRENT APPLICATION DATA:
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APPLICANT: Thomas Jr., Kenneth A.
TITLE OF INVENTION: VASCULAR ENDOTHELIAL CELL GROWTH FACTOR C
TITLE OF INVENTION: SUBUNIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                     APPLICANT: Bayne, Marvin L.
APPLICANT: Thomas Jr., Kenneth A.
TITLE OF INVENTION: VASCULAR ENDOTHELIAL CELL GROWTH FACTOR C
TITLE OF INVENTION: SUBUNIT
NUMBER OF SEQUENCES: 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER: 36,545
REFERENCE/DOCKET NUMBER: 18
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                        CORRESPONDENCE ADDRESS:
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STATE: New J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                    59 DEHPNEVSHIF 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: sin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE:
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                                                    STREET:
COUNTRY:
                                                                      ADDRESSEE:
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             : Rahway
E: New Jersey
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                                                  E: Merck & Co., Inc.
126 E. Lincoln Avenue
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Pred. No. 32;
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                                                                                                                                               US-09-173-300-34
                                                                                                                                                                                                            APPLICANT: Falco, Saverio Carl
APPLICANT: Hitz, William D.
APPLICANT: Kinney, Anthony J.
APPLICANT: Cahoon, Rebecca E.
APPLICANT: Cahoon, Rebecca E.
APPLICANT: Rafalski, J. Antoni
TITLE OF INVENTION: PLANT BRANCHED CHAIN AMINO ACID BIOSYNTHETIC ENZYMES
FILE REFERENCE: BB-1126
CURRENT FILING DATE: 1998-10-15
EARLIER APPLICATION NUMBER: 60/063,423
EARLIER APPLICATION NUMBER: 60/063,423
EARLIER FILING DATE: 1997 October 28
NUMBER OF SEQ ID NOS: 54
SOFTWARE: Microsoft Word Version 7.0A
SEQ ID NO 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 34, Applica
Patent No. 6451581
GENERAL INFORMATION:
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                                                                      Matches
                                                                                                       Query Match
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                ORGANISM: Escherichia coli
                                                                                                                                                                               TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                              LENGTH: 210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 36,545
REFERENCE/DOCKET NUMBER: 18
TELECOMUNICATION INFORMATION:
TELEPHONE: (908) 594-3905
TELEPAX: (908) 594-4720
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 08/124,259
FILING DATE: 20-SEP-1993
APPLICATION NUMBER: 07/676,436
FILING DATE: 28-MAR-1991
ATTORNEY/AGENT INFORMATION:
NAME: Hand, J. Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Microsoft Word 6
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: FILING DATE: 16-JAN CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        59 DEHPNEVSHIF 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 53.1%;
Local Similarity 54.5%;
les 6; Conservative
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36 DSHKGVVRHR 45
                                                                  Local Similarity
les 6; Conserv
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                                  1 DAHKSEVAHR 10
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                                                                      Conservative
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                                                                                       53.1%;
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Pred. No. 37;
                                                                      1; Mismatches
                                                                                       Score 34;
Pred. No.
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                                                                                         DB 4; Length 210; 50;
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Query Match
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US-08-357-533A-10
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US-08-158-735A-12
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                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: COOK, JONATHAN S.
APPLICANT: CORREA, PAUL E.
APPLICANT: KOENIG, BETH B.
APPLICANT: ROSENBAUM, JAN S.
APPLICANT: TING, JERRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (513) 627-0260 INFORMATION FOR SEQ ID NO: 1 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                   tent No.
COMPUTER READABLE FORM:
                                                                                                                                                     APPLICANT: JIN, DONALD F
APPLICANT: OPPERMANN, HERMANN
APPLICANT: KUBERASAMPATH, THANGAVEL
APPLICANT: KUBERASAMPATH, THANGAVEL
APPLICANT: SMART, JOHN E
TITLE OF INVENTION: NOVEL MORPHOGEN CELL SURFACE RECEPTOR
NUMBER OF SEQUENCES: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: 1; MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 34,804
REFERENCE/DOCKET NUMBER: 508
TELECOMMUNICATION INFORMATION:
TELEPHONE: (513) 627-2858
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/158,735A FILING DATE: 24-NOV-1993
                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: DNA SEQUENCE CODING FOR A BMP RECEPTOR NUMBER OF SEQUENCES: 22
                                                                                                                                                                                                                                                                                                                                                                                          122 DGHKPAISHR 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
              COUNTRY: US
ZIP: 01748
                                                              STREET: 45 SOUTH CITY: HOPKINTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino a
STRANDEDNESS:
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                                                                                                     ADDRESSEE: PAT
ADDRESSEE: INC
                                                                                                                                                                                                                                                                                                                                                                                                                            1 DAHKSEVAHR 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                45061
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                                                                                   45 SOUTH STREET
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11810 EAST MIAMI RIVER ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     323 amino acids
                                 USA
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                                                                                                                      PATENT ADMINISTRATOR, CREATIVE BIOMOLECULES
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Pred. No. 80;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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Query Match
Best Local Similarity
Matches 5; Conserve
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Patent No. 5861479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSER PADDRESS:
                APPLICATION NUMBER: US 08/357,533
FILING DATE: 16-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: KELLY, ROBIN D
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: CRP-073F
                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARB: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/357,533A
APPLICATION NUMBER: US/08/357,533A
                                                                                                  CLASSIFICATION: 435
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein FEATURE:
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LENGTH: 513 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: KELLY, ROBIN D
REGISTRATION NUMBER: 34,0
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                          STREET: 45 SOUTH CITY: HOPKINTON
                                                                                                                                                         FILING DATE:
                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                             ZIP: 01748
                                                                                                                                                                                                                                                                                                           COUNTRY:
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OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                             USA
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KUBERASAMPATH, THANGAVEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JIN,
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                                                                                                                                                                       US/08/459,009
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                     CRP-073FW
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Pred. No. 1.3e
2; Mismatches
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RESULT 50
US-08-300-584-2
; Sequence 2, Application US/08300584
; Patent No. 5885794
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LOCATION: 1..513
; OTHER INFORMATION: /note= "MOUSE ACTIVIN RECEPTOR"
US-08-459-009-10
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-300-584-2
                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:

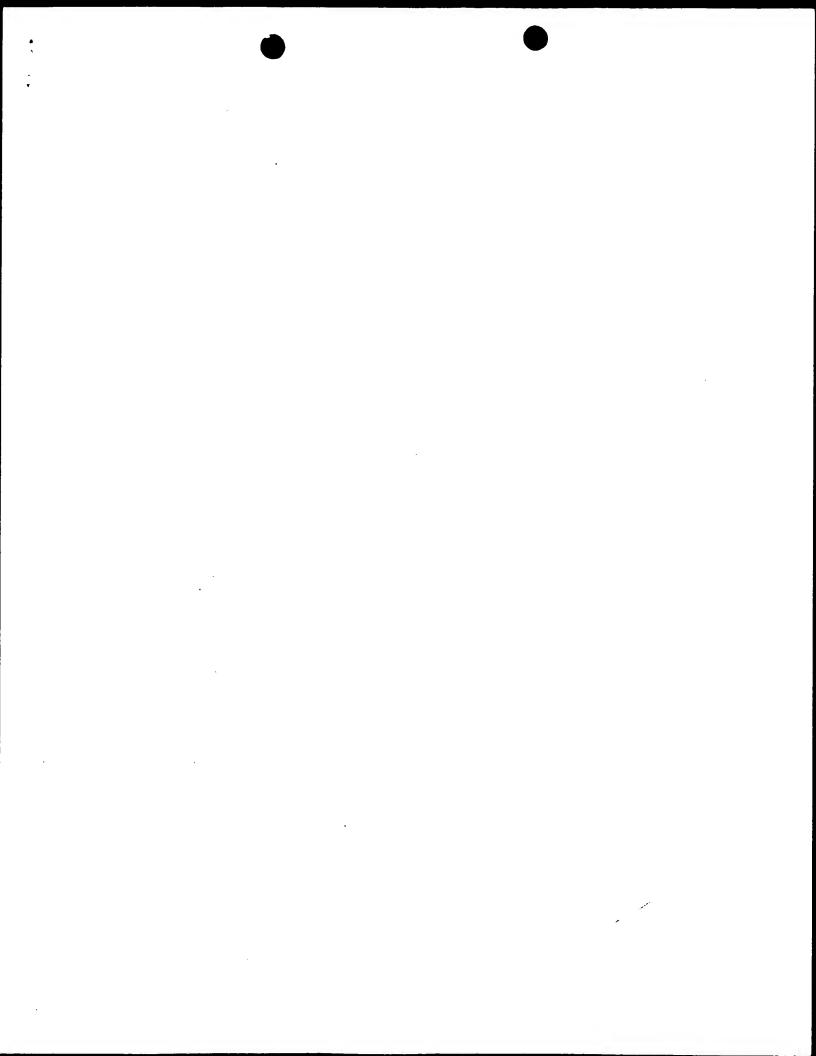
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNUMBER: US/08/300,584
FILING DATE: 02-SEP-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/880,220
FILING DATE: 08-MAY-1992
PRIOR APPLICATION DATA:
APPLICATION UNUMBER: US 07/773,229
FILING DATE: 09-OCT-1991
PRIOR APPLICATION DATA:
APPLICATION UNUMBER: US 07/698,709
FILING DATE: 10-MAY-1991
APPLICATION UNUMBER: US 07/698,709
FILING DATE: 10-MAY-1991
APPLICATION NUMBER: US 07/698,709
FILING DATE: 10-MAY-1991
APPLICATION DATE: US 07/698,709
FILING DATE: 10-MAY-1991
APPLICATION DATE: US 07/698,709
FILING DATE: 10-MAY-1991
APPLICATION DATE: US 07/698,709
FILING DATE: US 07/698,700
FILING DATE: US 07/698,700
FILING DATE: US 07/698,700
FILING DATE: US 07/698,700
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Best Local Similarity 50.0
Matches 5; Conservative
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TELEFAX: (508)-435-0992
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                            TELEFAX: 619-546-9392
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 513 amino acids
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APPLICANT: Vale, Wylle W.
TITLE OP INVENTION: CLONING AND RECOMBINANT PRODUCTION OF
TITLE OP INVENTION: RECEPTOR(S) OF THE ACTIVIN/TGF-BETA SUPERFAMILY
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSE: Pretty, Schroeder, Brueggemann & Clark
STREET: 444 South Flower Street, Suite 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: protein
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CITY: Los Angeles
STATE: CA
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TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
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Pred. No. 1.3e+02;
2; Mismatches 3
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Query Match
Best Local Similarity 50.0%; Pred. No. 1.3e+02;
Matches 5; Conservative 2; Mismatches 3; Indels 0; Gaps

Qy 1 DAHKSEVAHR 10
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Db 312 DGHKPAISHR 321

Search completed: April 11, 2003, 15:20:06

Job time: 16 secs
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Result
No.
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Maximum Match 100%
Listing first 75 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OM protein - protein search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Perfect score:
number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                           DB DB
                                                                                                                                                                                                                                         Score
                                                                                                                           39.5
                                                                                                                                                                                                                                                                             d. No. is the number of results predicted by chance to have a
re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                           seq length: 0
seq length: 2000000000
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64
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1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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DAHKSEVAHRFK 12
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A72691
S63392
T16270
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T02888
B70379
                                                                                           ABSHS
JC5838
JC5838
ABPGS
S57632
B61511
S29749
ABCHS
B82910
AH2378
S21171
JQ1317
AF3594
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A47391
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T30740
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ABRTS
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gephyrin homolog -
probable spermidin
probable membrane
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serum albumin - do
serum albumin prec
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ALIGNMENTS

ABHUS

serum albumin precursor [validated] - human

N;Alternate names: preproalbumin

N;Contains: kinetensin

C;Species: Homo sapiens (man)

C;Date: 29-Jul-1981 #sequence revision 31-Jan-1997 #text_change 17-Mar-2000

C;Date: 29-Jul-1981 #sequence . S.C.; Franke, A.E.; Houck, C.M.; Najarian, R.C.; Seeburg

R;Lawn, R.M.; Adelman, J.; Bock, S.C.; Franke, A.E.; Houck, C.M.; Najarian, R.C.; Seeburg

Nucleic Acids Res. 9, 6103-6114, 1981

A;Title: The sequence of human serum albumin cDNA and its expression in Escherichia coli.

A;Accession: A93743; MUID:82081882; PMID:6171778

A;Coss-references: EMBL:V00495; GB:J00078; GB:L00132; GB:L00133; NID:g28591; PIDN:CAA23:

A;Coss-references: EMBL:V00495; GB:L00132; GB:L00133; NID:g285

A; Molecule type: mRNA

protein F2

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A;Cross-references: EMBL:V00494; NID:g28589; PIDN:CAA23753.1; PID:g28590 R;Urano, Y.; Watanabe, K.; Sakai, M.; Tamaoki, T. J. Biol. Chem. 261, 3244-3251. J. Biol. Chem. 261, 3244-3251. A; Title: The human albumin gene. Characterization of the 5' and 3' flank: A;Reference number: I39427; MUID:86140099; PMID:2419329 A;Accession: I39427
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Molecule type: protein
A;Residues: 25-117,'EQ',120-154,'Q',156-193,'E',195-387,'H',389-390,'Y',392-393,'A',395-R;Roehr, U.; Spiteller, G.; Tripier, D.
Justus Liebigs Ann. Chem. 9, 881-884, 1988
A;Title: Isolation and structure elucidation of middle-molecular weight peptides from uz A;Reference number: S06422
A;Note: this paper is in German, with an English abstract
A;Accession: S06422
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A; Residues: 589-590; ALPRRVKNILLQVKLP' <MAD>
A; Residues: 589-590; ALPRRVKNILLQVKLP' <MAD>
A; Residues: GB:S70799; NID:g547231; PIDN:AAB31177.1; PID:g547232
A; Cross-references: GB:S70799; NID:g547231; PIDN:AAB31177.1; PID:g547232
A; NCte: this frame-shift variant is designated albumin Bazzano; four additional A; Note: this frame-shift variant is designated albumin Bazzano; four additional R; Menaya, J.; Parrilla, R.; Ayuso, M.S.
submitted to the EMBL Data Library, March 1995
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"Watkins, S.; Madison, J.; Galliano, M.; Minchiotti, L.; Putnam, F.W.
roc. Natl. Acad. Sci. U.S.A. 91, 2275-2279, 1994
"Title: A nucleotide insertion and frameshift cause analbuminemia in an Italian Reference number: I59286; MUID:94181575; PMID:8134387
"Accession: I59286
,;Molecule type: protein
,;Residues: 45-67;141-160;311-337,469-490;570-581 <FIN>
;;Residues: A: 59iteller, G.
;;Kausler, E.; Spiteller, G.
;;Causler, E.; Spiteller, G.
;;Causler, E.; Spiteller, G.
;;Causler, E.; Spiteller, G.
;Title: Bruchstuecke aus Albumin und beta(2)-Mikroglobulin -
,;Reference number: S17599; MUID:92126241; PMID:1772598
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;Residues: 282-290, 'KSRFDLQ' <WAT>
.Cross-references: GB:S69.192; NID:g546032; PIDN:AAB30282.1; PID:g546033
.Cross-references: GB:S69.192; NID:g546032; PIDN:AAB30282.1; PID:g546033
.Cross-references: GB:S69.192; NID:g546032; PIDN:AAB30282.1; PID:g546033
.Cross-reference: GB:S69.192; NID:g546032; PIDN:AAB30282.1; PID:g546033
.Cross-reference: Accession: J9313
.Cross-reference: Accession: I59313
.Cross-reference: I59313
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;Residues: 19-27 <LED>
;Residues: 19-27 <LED>
;Reloun, B.; Moravek, L.; Kostka, V.
EBS Lett 58, 134-137, 1975

Le: Complete amino acid sequence of human serum albumin.
Ference number: A91420; MUID:76187907; PMID:1225573
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, Residues: 1-120, 'G', 122-455 < MEN's
, Residues: 1-120, 'G', 122-455 < MEN's
, Residues: 1-120, 'G', 122-455 < MEN's
, Cross-references: EMBL: 1722961; NID: 9763428; PIDN: AAA64922.1; PID: 9763431
, Ledgerwood, E.C.; George, P.M.; Peach, R.J.; Brennan, S.O.
, Ledgerwood, E.C.; George, P.M.; Peach, R.J.; Brennan, S.O.
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Residues: 1-26 <URA>
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Biochem. Biophys. Res. Commun. 136, 983-988, 1986
A;Title: The amino acid sequence of kinetensin, a novel
A;Reference number: A03239; MUID:86242180; PMID:3087352
A;Accession: A03239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Title: Atomic structure and chemistry of human serum albumin. A;Reference number: A46756; MUID:93334427; PMID:1630489
A;Contents: annotation; X-ray crystallography, 2.8 angstroms R;Brown, J.R.; Shockley, P.; Behrens, P.O.
in The Chemistry and Physiology of the Human Plasma Proteins, E
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A; Residues: 255-263, 'E', 265-281 <MINI>
A; Residues: 255-263, 'E', 265-281 <MINI>
A; Note: this variant is designated albumin Herborn
A; Note: this variant is designated albumin Herborn
R; Minchiotti, L.; Galliano, M.; Stoppini, M.; Ferri, G.;
Biochim. Biophys. Acta 1119, 232-238, 1992
A; Title: Two alloalbumins with identical electrophoretic
A; Title: Two alloalbumins with identical electrophoretic
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A; Residues: 76-83, 'K', 85-106 <GAL3>
A; Residues: 76-83, 'K', 85-106 <GAL3>
A; Note: this variant is designated albumin Torino
A; Minchiotti, L.; Galliano, M.; Zapponi, M.C.; Tenni, R.
Bur. J. Biochem. 214, 437-444, 1993
A; Title: The structural characterization and bilirubin-binding
A; Reference number: S33298; MUID:93292504; PMID:8513793
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A;Residues: 166-173,'L' <MOG>
A;Residues: 166-173,'L' <MOG>
R;Galliano, M.; Minchiotti, L.; Porta, F.; Rossi, A.; Ferri, G.; Madison, J.;
R;Galliano, M.; Minchiotti, L. Porta, F.; Rossi, A.; Ferri, G.; Madison, J.;
Proc. Natl. Acad. Sci. U.S.A. 87, 8721-8725, 1990
A;Title: Mutations in genetic variants of human serum albumin found in Italy.
A;Reference number: A38255, MUID:91062352; PMID:2247440
A;Recession: C38255
                                                                                                                                                                                                                                                                                                                                                                    R;Saber, M.A.; Stockbauer, P.; Moravek, L.; Meloun, Collect. Czech. Chem. Commun. 42, 564-579, 1977 A;Title: Disulfide bonds in human serum albumin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Reference number: A94442
A; Contents: annotation; three-dimensional structure
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A;Residues: 354-356,'K',358-378 <MIN2>
A;Residues: 354-356,'K',358-378 designated albumin Sondrio; another variant A;Note: this variant is designated albumin Sondrio; another variant R;He, X.M.; Carter, D.C.
Nature 358, 209-215, 1992
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A;Residues: 82-105,'K',107-110 <GAL2>
A;Note: this variant is designated albumin Vibo Valentia
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A; Residues: 76-111 <GAL1:
A; Accession: B38255
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A; Reference number: A94408
A; Contents: annotation; bi
                                                                                                                                                   A;Title: Lysine residue 240 of human serum A;Reference number: A90299; MUID:78186630; A;Contents: annotation; bilirubin-binding s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Reference number: S21078; MUID:92190239; PMID:1347703
A;Accession: S21078
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                                                             A; Title: Serum albumin:
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                                                                                                                             R;Peters,
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                                                                                         in Albumin: Structure, Biosynthesis,
                                                                                                                                                                                                                                             J. 171, 453-459, 1978
                                                                                                                      T.; Reed, R.G.
                                                                                                                                                                                                                                                                                                          annotation; disulfide bonds
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   binding sites
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F;221-394/Domain: serum albumin repeat homology <SA2>
F;411-592/Domain: serum albumin repeat homology <SA3>
F;411-592/Domain: serum albumin repeat homology <SA3>
F;27/Binding site: copper (His) #status predicted
F;27/Binding site: copper (His) #status predicted
F;77-86,99-115,114-125,148-193,192-201,224-270,269-277,289-303,302-313,340-385,384-393,
F;214/Binding site: pyridoxal phosphate (Lys) (covalent) #status experimental
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C;Superfamily: serum albumin; serum albumin repeat homology
C;Superfamily: serum albumin; serum albumin repeat homology
C;Keywords: carrier protein; duplication; metal binding; phosphoprotein; plasma; pyridov
F;1-18/Domain: signal sequence #status predicted <SIG>
F;1-18/Domain: propeptide #status experimental <PNO>
F;1-24/Domain: propeptide #status experimental <PNO>
F;1-24/
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A;Note: the nonenzymatic transfer of an acetyl group from aspirin (acetylsalicyclic acia, R;Note: the nonenzymatic transfer of an acetyl group from aspirin (acetylsalicyclic acia, R;Nothney, J.P.; Fonda, M.L.; Feldhoff, R.C.
FEBS Lett. 298, 266-268, 1992
A;Title: Identification of Lys(190) as the primary binding site for pyridoxal 5'-phosph. A;Reference number: A56294; MUID:92183881; PMID:1544460
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R; Walker, J.E. FEBS Lett. 66, 173-175, 1976
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A;Reference number: A90028; MUID:83279982; PMID:6192711
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                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: GB:M90463; NID:g342294; PIDN:AAA36906.1; PID:g342295
A;Experimental source: liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Molecule type: mRNA; protein
A;Residues: 1-600 <WAT>
                                                                                                                                                                                                                                                                                                                                                                                                       A; Note: sequence extracted from NCBI backbone (NCBIN:128280,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Accession: A47391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Contents: B/B homozygote
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Watkins, S.; Sakamoto, Y.; Madison, J.; Davis, E.; Smith, D.G.; Dwulet, J.; Putnam, F. roc. Natl. Acad. Sci. U.S.A. 90, 2409-2413, 1993
Title: cDNA and protein sequence of polymorphic macaque albumins that differ in bilirus Reference number: A47391; MUID:93211971; PMID:8460152
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                                                                                                                                                                                              Superfamily: serum albumin; serum albumin repeat homology;21-194/Domain: serum albumin repeat homology <8A1-;213-386/Domain: serum albumin repeat homology <8A2-;405-584/Domain: serum albumin repeat homology <8A3-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Status: preliminary
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Species: Macaca mulatta (rhesus macaque)
Date: 21-Jan-1994 #sequence_revision 18-Nov-1994 #text_change 20-Aug-1999
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                                               Local Similarity
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                                               93.8%;
91.7%;
Score 60; DB 2;
Pred. No. 0.0041;
0; Mismatches
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Pred. No. 0.00077;
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                                                                                                    DB 2; Length 600;
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A;Accession: A;Accession A;Acc
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A;Residues: 1-32 <MAG>
R;Hsieh, J.C.; Lin, F.P.; Tam, M.F.
Anal. Biochem. 170, 1-8, 1988
A,Title: Electroblotting onto glass-fiber filter from an analytical isoelectrofocusing g@a,Title: Electroblotting onto glass-fiber filter from an analytical isoelectrofocusing g@a,Reference number: A60808; MUID:88267456; PMID:3389500
A;Accession: B60808
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Eur. J. Biochem. 98, 477-485, 1979

A;Title: Biosynthesis of bovine plasma proteins in a cell-free system. A;Reference number: A91258; MUID:80024278; PMID:488109

A;Accession: A91258
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R;Hirayama, K.; Akashi, S.; Furuya, M.; Fukuhara, K.
Biochem. Biophys. Res. Commun. 173, 639-646, 1990
A;Title: Rapid confirmation and revision of the primary structure of bovine serum albumin A;Reference number: A36401; MUID:91083649; PMID:2260975
A;Accession: A36401
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C;Date: 24-Apr-1984 #sequence revision 30-Sep-1993 #text change 18-Aug-2000 C;Dates: 24-Apr-1984 #sequence revision 30-Sep-1993 #text change 18-Aug-2000 C;Accession: A38885; A36401; Ā91258; B66808; S10780; D45800; A26693; A90309; R;Holowachuk, E.W.; Stoltenborg, J.K.; Reed, R.G.; Peters Jr., T. submitted to the EMBL Data Library, August 1991 A;Description: Bovine serum albumin: cDNA sequence and expression.
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A;Residues: 25-41, 'H', 43-57,59-64 <STR>
A;Residues: 25-41, 'H', 43-57,59-64 <STR>
R;Carraway, R.E.; Cochrante, D.E.; Boucher, W.; Mitra, S.P.

J. Immunol. 143, 1680-1684, 1989
A;Title: Structures of histamine-releasing peptides formed by the action of acid protease
A;Reference number: A45800; MUID:89341406; PMID:2474609
A;Accession: D45800
A;Accession: D45800
A;Molecule type: protein
A;Residues: 163-172 <CARP
C;Carraway, R.E.; Mitra, S.P.; Cochrane, D.E.
J. Biol. Chem. 262, 5968-5973, 1987
A;Title: Structure of a biologically active neurotensin-related peptide obtained from per
A;Accession: A26693; MUID:87194805; PMID:2437111
A;Accession: A26693
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A;Title: Tooth 'enamelins' identified mainly as serum proteins. Major 'enamelin' is album A;Reference number: S10780; MUID:90336641; PMID:2379503
A;Accession: S10780
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Residues: 1-607 < HOL>
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R;Strawich, E.; Glimcher, M.J.
                                                                                                                                                                                                                                                                                                                               A; Reference number: A90309; A; Accession: A90309
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                                                A; Title: Structure of
                                                                                                                                                            R; Brown, J.R
                                                                                                                                                                                                            A; Molecule type: protein
A; Residues: 402-433 < REE:
                                                                                                      Fed. Proc. 34, 591,
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number:
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A91458
                                                                                                            1975
                                                      bovine
                                                      serum albumin
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A;Molecule type: protein
A;Residues: 1-38 <STR>
A;Residues: 3-31 <STR>
R;Isemura, 5; Ikenaka, T.
J. Biochem. 83, 35-48, 1978
A;Title: Amino acid sequences of fragments I and II obtained by cyanogen bromide cleavag A;Reference number: A91946; MUID:78109429; PMID:564345
A;Accession: A91946
A;Residues: 25-222 <ISL)
R;Isemura, S.; Ikenaka,
                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: GB:V01222; GB:J00698; NID:g55627; PIDN:CAA24532.1; PID:g55628 R;Strauss, A.W.; Bennett, C.D.; Donohue, A.M.; Rodkey, J.A.; Alberts, A.W. J. Biol. Chem. 252, 6846-6855, 1977 A;Title: Rat liver pre-proalbumin: complete amino acid sequence of the pre-piece. Analys A;Reference number: A92211; MUID:77249657; PMID:893447
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N;Alternate names: preproalbumin (Norway rat)
C;Speciles: Rattus norvegicus (Norway rat)
C;Date: 31-May-1979 #sequence revision 31-May-1979 #text change 22-Jun-1999
C;Accession: A93872; A92211; A91946; A91940; C45800; 157621; A03233
Yegent, T.D.; Yang, M.; Bonner, J.
Natl: Acad. Sci. U.S.A. 78, 243-246, 1981
A;Title: Nucleotide sequence of cloned rat serum albumin messenger RNA. A;Reference number: A93872; MUID:81223722; PMID:7017712
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A;Title: Preparation and characterization of novel substrates of insulin proteinase A;Title: Preparation and characterization of novel substrates of insulin proteinase A;Reference number: S55232; MUID:95031935; PMID:7945219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             serum albumin precursor - rat
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A;Residues: 25-41,'H',43-117,'EQ',120-179,181-189,'E',191-194,'A',196-213,'T',215-288,'
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                                            Molecule type: protein
Residues: 25-222 <IS1>
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Residues: 1-608 <SAR>
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;412-591/Domain: serum albumin repeat homology <SA3>
;27/Binding site: copper (His) #status predicted
;77-86,99-115,114-125,147-192,191-200,223-269,268-276,288-302,301-312,339-384,383-392,
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;Residues: 190-195 <BR2>
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sidues: 529-536;569-572 <WER>
sidues: 529-536;569-572 <WER>
partiments: serum albumin; serum albumin repeat homology
(keywords: carrier protein; copper binding; duplication; p
1-18/Domain: signal sequence #status experimental <SIG>
19-24/Domain: propeptide #status experimental <PRO>
25-607/Product: serum albumin #status experimental <MPT>
259-201/Domain: serum albumin repeat homology <SA1>
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83.3%;
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Pred. No. 0.0064;
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F;220-393/Domain: serum albumin repeat homology <SA2>
F;412-591/Domain: serum albumin repeat homology <SA3>
F;27/Binding site: copper (His) #status predicted
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A;Title: Determinants of rat albumin promoter tissue specificity analyzed by an improved
A;Reference number: I57621; MUID:87286876; PMID:3475566
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R.Aoyagi, Y; Ikenaka, T; Ichida, F.

Cancer Res. 38, 3483-3486, 1978
A; Title: Copper(III-binding ability of human alpha-fetoprotein.

A; Reference number: A99758; MUID: 79001617; PMID: 80265
A; Contents: annotation; copper binding
A; Carraway, R.E.; Cochrane, D.E.; Boucher, W.; Mitra, S.P.
                                                                                                          F;25-607/Product: serum albumin #status predicted <MAT>F;29-201/Domain: serum albumin repeat homology <SA1>
                                                                                                                                                                                    F;19-24/Domain: signal sequence #status predicted <SIG>F;19-24/Domain: propeptide #status predicted <PRO>
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Nucletc Acids Res. 17, 10495, 1989
A;Title: Nucleotide and deduced andno acid sequence of sheep serum albumin A;Reference number: S06936; MUID:90098888; PMID:2602160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                serum albumin precursor - sheep
C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 22-Jun-1999
C;Accession: S06936
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J. Immunol. 143, 1680-1684, 1989
A,Title: Structures of histamine-releasing peptides formed by the action of acid protease A,Reference number: A45800; MUID:89341406; PMID:2474609
A,Accession: C45800
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A; Residues: 1-607 < BRO>
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F;413-592/Domain: serum albumin repeat homology <8A3>
F;27/Binding site: copper (His) #status experimental
F;77-86,99-115,114-125,148-193,192-201,224-270,269-277,289-303,302-313,340-385,384-393,41
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A; Residues: 1-5 < RES>
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A; Residues: 223-288;572-608 <IS2>
A; Note: 262-Leu was also found
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A;Accession: A91940
                                                                                                                                                                                                                                                               Superfamily: serum albumin; serum albumin repeat homology Roywords: carrier protein; duplication; metal binding; plasma
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Local Similarity 83.3%;
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Pred. No. 0.0064;
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F;25-607/Product: Berum albumin #status predicted <MAT>
F;29-201/Domain: Berum albumin repeat homology <SA1>
F;290-393/Domain: Berum albumin repeat homology <SA2>
F;200-393/Domain: Berum albumin repeat homology <SA3>
F;412-591/Domain: Berum albumin repeat homology <SA3>
F;412-591/Domain: Berum albumin repeat homology <SA3>
F;27/Binding Bite: Copper (His) #status predicted
F;77-86,99-115,114-125,147-192,191-200,223-269,268-276,288-302,301-312,339-384,383-392,
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DNA Res. 4, 351-354, 1997
A;Title: Sequencing of cDNA encoding serum albumin and its extrahepatic synthesis in the A;Reference number: JC5838; MUID:98116663; PMID:9455485
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C;Comment: Serum albumin is synthesized in the liver as preproalbumin. It binds copper, teroid hormones (weak bonds with these hormones promote their transfer across the membra
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F;263/Binding site: bilirubin (Lys) #status predicted
                                                                                                                                                                                                                                                                                                                                                  A; Molecule type: mRNA
A; Residues: 1-609 <70S
A; Cross-references: DDBJ:AB006197; NID:g2317277; PIDN:BAA21765.1; PID:g2317278
A; Experimental source: liver
C; Superfamily: serum albumin; serum albumin repeat homology
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C;Species: Equus caballus (domestic horse)
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A; Residues: 1-607 < HOA>
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                                                                                                                                                                                                                                                                                                                     ;222-395/Domain: serum albumin repeat homology <SA2>
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;Date: 05-Mar-1998 #sequence_revision 13-Mar-1998 #text_change 19-May-2000
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Lle: X-ray and primary structure of horse serum albumin (Equus caballus) at 0.27-nm Reference number: $34053; MUID:93345495; PMID:8344282
                                                                                                                                       Matches
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(es 9; Conservation
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Pred. No. 0.053;
1; Mismatches
                                                                                                                                       2; Mismatches
                                                                                                                                                                                   Score 54; DB 2; Length 609; Pred. No. 0.053;
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C;Species: Sus scrofa domestica (domestic pig)
C;Date: 31-Dec-1993 #sequence revision 31-Dec-1993 #text_change 22-Jun-1999
C;Accession: S01382, A61006
R;Weinstock, J; Baldwin, G.S.
Nucleic Acids Res. 16, 9045, 1988
A;Title: Nucleotide sequence of porcine liver albumin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Superfamily: serum albumin; serum albumin repeat homology
C;Keywords: carrier protein; duplication; metal binding; plasma
F;11-5(Domain: signal sequence (fragment) #status predicted <SIG>
F;17-22/Domain: propeptide #status predicted <PRO>
F;23-065/Product: serum albumin #status predicted <MAT>
F;23-07-199/Domain: serum albumin repeat homology <SA1>
F;218-391/Domain: serum albumin repeat homology <SA2>
F;410-589/Domain: serum albumin repeat homology <SA3>
F;410-589/Domain: serum albumin repeat homology <SA3-
F;410-589/Domain: s
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A; Residues: 23-51, 'X', '53-54; 'XXXGY', 146, 'E', 148, 'E', 150-151, 'XVN', 155 <LIM>A; Residues: 23-51, 'X', '53-54; 'XXXGY', 146, 'E', 148, 'E', 150-151, 'XVN', 155 <LIM>A; Rexperimental source: dental enamel
A; Note: albumin and other serum proteins are also found in bone
A; Note: albumin and other serum proteins are also found in bone
A; Note: albumin and other serum proteins are also found in bone
A; Note: albumin and other serum proteins are also found in bone
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A; Note: albumin and other serum proteins are also found in bone
A; Note: albumin and other serum proteins are also found in bone
A; Note: albumin and other serum proteins are also found in bone
A; Note: albumin albumin is synthesized in the liver as preproalbumin. It because albumin albumin is synthesized in the liver as preproalbumin.
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J. Bone Miner. Res. 4, 235-241, 1989
A;Title: Serum albumin and its acid hydrolysis peptides
A;Reference number: A61006; MUID:89269769; PMID:2728927
A;Accession: A61006
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                                 C;Keywords: liver; plasma
F;1-18/Domain: signal sequence #status predicted <SIG>F;19-24/Domain: propeptide #status predicted <PRP>F;25-608/Product: serum albumin #status predicted <MAT>F;29-202/Domain: serum albumin repeat homology <SAl>
                                                                                                                                                                                                                                                                     C;Comment: This protein is the major protein component in plasma. ein has 35 conserved cysteine residues.
C;Superfamily: serum albumin; serum albumin repeat homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   serum albumin precursor - cat (;Species: Felis silvestris catus (domestic cat) C;Species: Felis silvestris catus (domestic cat) C;Date: 19-Oct-1995 #sequence revision 03-Nov-1995 #text_change 20-Aug-1999 C;Accession: JC4660; S57632
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                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: EMBL:X84842; NID:g886484; PIDN:CAA59279.1; PID:g886485
A;Experimental source: liver
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A; Residues: 1-608 <HI2>
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A; Reference number: JC4660; MUID: 96194824;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Hilger, C.; Grigioni, F.; Hentges, F. Gene 169, 295-296, 1996
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         F;221-394/Domain: serum albumin repeat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 DAHKSEVAHRFK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AHKSEIAHRYK 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           translation not shown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       82.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 53; DB 1; Length 605
Pred. No. 0.081;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PMID:8647469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (Felis domesticus)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          serum albumin
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a multi

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serum albumin precursor - chicken C;Species: Gallus gallus (chicken) C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 22-Jun-1999 C;Accession: S15571; A05078; A13451 R;Cassady, A.I.; Salklld, C.K.; Baverstock, P.; Wallace, J.C. submitted to the EMBL Data Library, July 1991 A;Reference number: S15571 A;Accession: S15571 A;Accession: S15571
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Residues: 1-30
C; Superfamily: s
C; Keywords: milk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  serum albumin, milk-derived - Australian echidna (fragment)
C;Species: Tachyglossus aculeatus (Australian echidna)
C;Date: 09-Oct-1994 #sequence_revision 15-Oct-1994 #text_change 11-May-2000
C;Accession: B61511
                                                                                                                                                                          ABCHS
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J. Biol. Chem. 249, 5872-5877, 1974

A;Title: Isolation, amino acid sequence and copper(II)-binding

A;Reference number: S29749; MUID:75011422; PMID:4414013
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Species: Canis lupus familiaris (dog)
;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 07-May-1999;Accession: S29749
                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Superfamily: serum albumin;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tatus: preliminary
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). Biochem. Physiol. B 9, 99-118, 1991
ttle: Some monortreme milk "whey" and blood proteins.
ference number: A61511; MUID:92070088; PMID:1959333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cession: B61511
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les 8; Conser
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1 EAYKSEIAHRY 11
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                     70.3%;
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72.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 serum albumin repeat homology
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                                                                                                                                                                                                                                                                                                                                       Score 45;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 50; DB 2
Pred. No. 0.29;
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 46; DB 2;
Pred. No. 0.068;
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                                                                                                                                                                                                                                                                                                                                       0.083;
                                                                                                                                                                                                                                                                                                                                                          DB 2;
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C;Comment: Serum albumin is synthesized in the liver as preproalbumin. mones (weak bonds with these hormones promote their transfer across the C;Superfamily: serum albumin; serum albumin repeat homology C;Keywords: carrier protein; duplication; metal binding; plasma F;1-18/Domain: signal sequence #status predicted <SIG> F;1-26/Domain: propeptide #status predicted <PRO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hypothetical protein UU287 [imported] - Ureaplasma urealyticum
C;Species: Ureaplasma urealyticum
C;Date: 18-Aug-2000 #sequence_revision 20-Aug-2000 #text_change 20-Aug-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: protein
A; Residues: 19-23, 'M', 25-30 <ROS>
C; Comment: Serum albumin is synth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Reference number: Al3451; MUID:78019943; PMID:911327
A;Accession: Al3451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Residues: 1-615 <CAS>
A;Cross-references: EMBL:X60688; NID:g63747; PIDN:CAAA3098.1; PID:g63748
A;Cross-references: EMBL:X60688; NiD:g63747; PIDN:CAAA3098.1; PID:g63748
R;Hache, R.J.G.; Wiskocil, R.; Vasa, M.; Roy, R.N.; Lau, P.C.K.; Deeley, R.G.
J. Biol. Chem. 258, 4556-4564, 1983
A;Title: The 5' noncoding and flanking regions of the avian very low density apolipoprote
A;Reference number: A05078; MUID:83161037; PMID:6187737
밁
                                                                                                                                                                                                                                                                              A;Residues: Î-201 <GLA>
A;Cross-references: GB:AE002125; GB:AF222894; NID:g6899253; PIDN:AAF30696.1; GSPDB:GN001;
A;Experimental source: serovar 3; biovar 1
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Accession: E82910
R;Glass, J.I.; Lefkowitz, E.J.; Glass, J.S.; Heiner,
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F;417-596/Domain: serum albumin repeat homology <8A3>
F;417-596/Domain: serum albumin repeat homology <8A3>
F;30/Binding site: copper (His) #stratus predicted
F;80-89,102-118,117-128,152-197,196-205,228-274,273-281,293-307,306-317,344-389,388-397,4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Rosen, A.M.; Geller, D.M.
Biochem. Biophys. Res. Commun. 78, 1060-1066, 1977
A;Title: Chicken microsomal albumin: amino termina
                                                                                                                                                                                                                         A;Gene: UU287
A;Genetic code: SGC3
                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                      A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Accession: E82910
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Reference number: A82870
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Description: The complete sequence of Ureaplasma urealyticum: Alternate views of a min:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      submitted to GenBank, February 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E82910
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               F;27-613/Product: serum albumin #status predicted <MAT>F;32-206/Domain: serum albumin repeat homology <SA1>
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A; Residues: 1-28 <HAC>
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79
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                                                                                                                                       Local Similarity
                                                     3 HKS-----EVAHRFK 12
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  HKSSSLYLTNDDEVAHRFK
                                                                                                            10;
                                                                                                            Conservative
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                                                                                                                                       61.7%;
52.6%;
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  97
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                                                                                                                                       Score 39.5;
Pred. No. 7.
                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C.R.; Chen, E.Y.; Cassell,
                                                                                                                                                                  Length 201;
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                                                                                                            Gaps
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hypothetical protein asr4584 [imported] - Nostoc sp. C; Species: Nostoc sp.

A; Note: Nostoc

вp.

strain PCC 7120 is a

synonym of Anabaena

PCC 7120

(strain PCC 7120)

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A;Cross-references: GB:S70930; NID:g240781; PIDN:AAB20638.1; PID:g240782
C;Superfamily: activin receptor II; protein kinase homology
C;Keywords: ATP; glycoprotein; serine/threonine-specific protein kinase; transmembrane
F;1-20/Domain: signal sequence #status predicted <SIG>
F;21-514/Product: activin receptor #status predicted <ACT>
F;317-162/Domain: transmembrane #status predicted <TRA>
F;191-487/Domain: protein kinase homology <XIN>
F;46,67,88,214,334/Binding site: carbohydrate (Asn) (covalent) #status predicted
                                                                                                                                                                                                                                                                                                  C;Accession: JQ1317
R;KONDO, M.; Tashiro, K.; Fujii, G.; Asano, M.; Miyoshi, R.; Yamada, R.; Muramatsu, Biochem. Biophys. Res. Commun. 181, 684-690, 1991
Biochem. Biophys. Res. Commun. 181, 684-690, 1991
A;Title: Activin receptor mRNA is expressed early in Xenopus embryogenesis and the A;Reference number: JQ1317; MUID:92095974; PMID:1661587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Superfamily: activin receptor 11; protein vini
C;Keywords: ATP
F;189-485/Domain: protein kinase homology <KIN>
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A;Title: Complete Genomic Sequence of the Filamentous Nitrogen-fixing Cyanobacterium An: A;Reference number: AB1807; MUID:21595285; PMID:11759840

A;Accession: AH2378
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A;Residues: 1-512 <NIS>
C;Superfamily: activin receptor
C;Keywords: ATP
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FEBS Lett. 303, 81-84, 1992
A;Title: Multiple genes for Xenopus activin receptor expressed during early embryogenesi A;Reference number: S21171; MUID:92275088; PMID:1317302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             activin receptor STK9 - African clawed frog)
C;Species: Xenopus laevis (African clawed frog)
C;Species: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 28-Feb-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                               activin receptor precursor - African clawed frog
C;Species: Xenopus laevis (African clawed frog)
C;Date: 31-Mar-1992 #sequence_revision 31-Mar-1992 #text_change 18-Jun-1999
C;Accession: JQ1317
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A;Experimental source: strain PCC 7120
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A; Residues: 1-93 < KUR>
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A; Residues: 1-514 < KON>
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hypothetical protein 138R - Molluscum contagiosum virus 1 N;Alternate names: MC138R C;Species: Molluscum contagiosum virus 1 C;Date: 05-Nov-1999 #sequence_revision 05-Nov-1999 #text c
                                                                                                               RESULT
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R;Delvecchio, V.G.; Kapatral, V.; Redkar, R.J.; Patra, G.; Mujer, C.; Los, T.; Ivanova, P.; Mazur, M.; Goltsman, E.; Selkov, E.; Elzer, P.H.; Hagius, S.; O'Callaghan, D.; Letessc Proc. Natl. Acad. Sci. U.S.A. 99, 443-448, 2002
A;Title: The genome sequence of the facultative intracellular pathogen Brucella melitensi A;Reference number: AD3252; PMID:11756688
A;Accession: AF3594
                                                                                                                                                                                                                                                                                                              K;Michalowski, C.B.; Bohnert, H.J.
submitted to the EMBL Data Library, March 1998
A;Description: An expressed sequence tag for sucrose synthase
A;Reference number: Z17473
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                                                                                                                                                                                                                                                                                                                                                                                                                                   sucrose synthase (EC 2.4.1.13) - common ice plant (fragment) C;Species: Mesembryanthemum crystallinum (common ice plant) C;Date: 23-Jul-1999 #sequence_revision 23-Jul-1999 #text_cha
                                                                                                                                                                     A;Cross-references: EMBL:AF054446; NID:g3064040; PID:g3064041
C;Superfamily: sucrose synthase; sucrose/sucrose-phosphate synthase homology
C;Keywords: glycosyltransferase; hexosyltransferase
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A; Residues: 1-67 < MIC>
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A; Residues: 1-489 < KUR>
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Pred. No.
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m contagrosum virus 1
#sequence_revision 05-Nov-1999 #text_change 21-Jul-2000

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probable molybdopterin synthase large subunit [imported] - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 02-Feb-2001
C;Accession: B84870
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Best Local Similarity
Thes 7; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RiStover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, P.; Hickey, M.J.; Bradman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, A.; Larbig, K.; Lim, Lory, S.; Olson, M.V.
Nature 406, 959-964, 2000
A;Title: Complete genome sequence of Pseudomonas aeruginosa PA01, an opportunistic patho A;Reference number: A82950; MUID:20437337; PMID:10984043
A;Accession: F83532
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R;Senkevich, T.G.; Bugert, J.J.; Sisler, J.R.; Koonin, E.V.; Darai, Science 273, 813-816, 1996
A;Gene: At2g43760
A;Map position: 2
                                                        A;Cross-ref
C;Genetics:
                                                                                                           A; Molecule type: DNA
A; Residues: 1-198 <STO>
                                                                                                                                                                                          A;Title: Sequence and analysis of chromosome 2 of the plant Arabidopsis thaliana. A;Reference number: A84420; MUID:20083487; PMID:10617197 A;Accession: B84870
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C;Species: Pseudomonas aeruginosa
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C;Superfamily: variola major virus 6R protein
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A; Residues: 1-117 <SEN>
                                                                                                                                                                 A;Status: preliminary
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                                                                                    Cross-references:
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Residues: 1-153 <STO>
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                                                                                  GB:AE002093; NID:g2281094; PIDN:AAB64030.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        56.2%;
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77.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 36;
Pred. No.
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Pred. No. 19;
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                                                                                    GSPDB:GN00139
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                            T42407
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gephyrin homolog
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A;Gene: TK
C;Superfamily: Escherichia coli thymidine kinase
C;Keywords: ATP; DNA biosynthesis; phosphotransferase
                                                                                                                                                                                                                                                                                                                                                                                         C;Species: Aquifex aeolicus
C;Date: 08-May-1998 #sequence_revision 08-May-1998 #text_change 11-Jun-1999
C;Accession: B70379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Ullah, H.; Robertson, N.; Fites, R.C. submitted to the EMBL Data Library, May 1998 A;Description: Plant thymidine kinase.
                                                                                                                A;Gene: dapB
C;Superfamily: dihydrodipicolinate reductase
                                                                                                                                                          A;Experimental source: strain C;Genetics:
                                                                                                                                                                                                                                                                               A;Title: The complete genome of the hyperthermophilic bacterium Aquifex aeolicus A;Reference number: A70300; MUID:98196666; PMID:9537320 A;Accession: B70379
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A;Experimental source: cultivar Nipponbare
C;Genetics:
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                                                                                                                                                                                                A; Cross-references: GB: AE000713; NID: g2983424; PIDN: AAC07008.1;
                                                                                                                                                                                                                    A;Residues: 1-265 <AQF>
                                                                                                                                                                                                                                         A; Molecule type: DNA
                                                                                                                                                                                                                                                        A; Status: preliminary; nucleic acid sequence not shown; translation not shown
                                                                                                                                                                                                                                                                                                                                           Nature 392, 353-358, 1998
                                                                                                                                                                                                                                                                                                                                                                                 R;Deckert, G.; Warren, P.V.; Gaasterland, T.; Young, W.G.; Lenox, A.L.; Graham,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              dihydrodipicolinate reductase - Aquifex aeolicus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Residues: 1-212 <ULL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Molecule type: mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Reference number: Z14763
A; Accession: T02888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Species: Oryza sativa (rice)
C;Date: 24-Mar-1999 #sequence_revision 24-Mar-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     thymidine kinase (EC 2.7.1.21) [similarity] - rice
                                       Matches
                                                          Query Match
Best Local
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Best Local Similarity
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1 DAHKSEVAHRFK 12
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                                     Similarity 7; Conserv
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                                     Conservative
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58.3%;
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70.0%;
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                                                        Score 36;
Pred. No.
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Pred. No.
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                                         Mismatches
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                                                                             Length 265
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151 DAEIMEIHHRFK 162

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Caenorhabditis elegans

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C;Species: Caenorhabditis elegans
C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 17-Mar-2000
C;Accession: T42407
R;Moss, E.G.; Johnstone, J.; Lee, R.; Ambros, V.
submitted to the EMBL Data Library, January 1998
A;Reference number: Z22161
A;Accession: T42407
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Rolecule type: mRNA
A;Residues: 1-391 < MOSS  
A;Cross-references: EMBL:AF042069; PIDN:AAB97841.1
C;Gene: lin-46
C;Superfamily: molybdenum cofactor biosynthesis protein moeA-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R; Kawarabayasi, Y.; Hino, Y.; Horikawa, H.; Yamazaki, S.; Haikawa, Y.; Jin-no, K.; Takalawa, H.; Takaniya, M.; Masuda, S.; Funahashi, T.; Tanaka, T.; Kudoh, Y.; Yamazaki, J.; PDNA Res. 6, 83-101, 1999
A; Title: Complete genome sequence of an aerobic hyper-thermophilic Crenarchaeon, Aeropyr A; Reference number: A72450; MUID:99310339; PMID:10382966
                                                                              A;Molecule type: DNA
A;Residues: 1-719 <DUE>
A;Cross-references: EMBL:271675; NID:g1302583; PIDN:CAA96342.1;
A;Experimental source: strain S288C
                                                                                                                                                                                                                                                                                 C;Species: Saccharomyces cerevisiae
C;Date: 27-Apr-1996 #sequence_revis:
C;Accession: S63392
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A;Molecule type: DNA
A;Residues: 1-407 <KAW>
A;Cross-references: DDBJ:AP000060; NID:g5104188; PIDN:BAA79929.1; PID:d1043715; PID:g510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        probable spermidine/putrescine-binding periplasmic protein APE0945 - Aeropyrum pernix C;Species: Aeropyrum pernix C;Date: 20-Aug-1999 #sequence_revision 20-Aug-1999 #cext_change 08-Dec-2000 C;Accession: A72691
                                                                                                                                                                                                            R;Duesterhoeft, A.; Floeth, M.; Fritz, C.; Heuss-Neitzel, D.; Hilbert, H.; Moestl, submitted to the Protein Sequence Database, April 1996
A;Reference number: S62944
                                                                                                                                                                                                                                                                                                                                                        probable membrane protein YNR060\mbox{w} - yeast (Saccharomyces cerevisiae) N;Alternate names: hypothetical protein N3518
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Gene: APE0945
C;Superfamily: Escherichia coli spermidine/putrescine-binding protein
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                                              A;Gene: SGD:FRE4
                                                                                                                                                                                        A;Accession: S63392
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                        Cross-references: SGD:S0005343; MIPS:YNR060w
                                                                                                                                                                                                                                                                                                       Date: 27-Apr-1996 #sequence_revision 03-May-1996 #text_change 20-Jun-2000
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position:
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Local Similarity 63.6%;
les 7; Conservation
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Pred. No. 68;
2; Mismatches
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                                                 Matches
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A; Description: The sequence of A; Reference number: Z18487 A; Accession: T16270
                                                                                                                                                                                                                                                                                                                                                                                                                                             hypothetical protein F35D11.11 - Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Date: 20-Sep-1999 #sequence_revision 20-Sep-1999 #text_change 20-Sep-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Superfamily: ferric reductase FRE2
C;Keywords: transmembrane protein
F;1-17/Domain: transmembrane #status predicted <TMl>
                                                                                                                                                                                                  A;Molecule type: DNA
A;Residues: 1-1827 <FUL>
A;Cross-references: EMBL:U29381; NID:g868214; PID:g868224; PIDN:AAA68757.1; CESP:F35D11.1
A;Experimental source: strain Bristol N2
                                                                                                                                                                                                                                                                                                                                                       submitted to the EMBL Data Library, June 1995 A;Description: The sequence of C. elegans cosmid F35D11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         F;374-390/Domain: transmembrane #status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              F;350-366/Domain:
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                                                                                                                                 A; Introns: 76/2; 131/3; 159/3; 185/3; 221/3; 253/3; 320/1; 869/3; 1133/3; 1205/2;
                                                                                                                                                           A;Gene: CESP:F35D11.11
                                                                                                                                                                              C; Genetics:
                                                                                                                                                                                                                                                                                        A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                               C; Accession:
                                                                                     Query Match
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                                                                   Local
3 HKSEVAHRFK 12
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                                            Similarity 6; Conserv
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                                              Conservative
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75.0%;
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                                            2; Mismatches
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                                                                Score 36; DB 2;
Pred. No. 3.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 36; DB 2;
Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      predicted <TM6>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 719;
                                              2; Indels
                                                                                     Length 1827
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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A;Authors: Utterback, T.; Cotton, M.D.; Spriggs, T.; Artiach, P.; Kaine, B.P.; Sykes, Smith, H.O.; Woses, C.R.; Venter, D.G., Tritle: The complete genome sequence of the hyperthermophilic, sulfate-reducing archa, Reference number: A69250; MUID:98049343; PMID:9389475
A;Accession: G69489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LSU ribosomal protein L22P (rpl22P) homolog - Archaeoglobus fulgidus C;Species: Archaeoglobus fulgidus C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 13-Aug-1999
                                                                                                                                                                                                                                                                                                                                                                                Glodek, A., Zhou, L., Ov.
Nature 390, 364-370, 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Accession:
R;Klenk, H.P
                                                                                                       A;Cross-references: GB:AE000971; GB:AC0;Superfamily: rat ribosomal protein
                                                                                                                                                           A; Residues: 1-155 < KLE>
                                                                                                                                                                                     A; Molecule type: DNA
                                                                                                                                                                                                               A;Status: preliminary; nucleic acid sequence not shown; translation
                                                                                                                                                                                                                                                                                                                                                                                                          ;Klenk, H.P.; Clayton, R.A.; Tomb, J.F.; White, O.; Nelson, K.E.; Ketchum, K.A.; Dodson, Fleischmann, R.D.; Quackenbush, J.; Lee, N.H.; Sutton, G.G.; Gill, S.; Kirkness, E.F.; Glodek, A.; Zhou, L.; Overbeek, R.; Gocayne, J.D.; Weidman, J.F.; McDonald, L.
                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              G69489
                         Similarity
Conservative
                                                                                                          GB:AE000971; GB:AE000782; NID:g2689294; ribosomal protein L17
                       54.7%;
75.0%;
                       Score 35;
Pred. No.
  Mismatches
                    DB 40;
                                                   2
                                                   Length 155
Indels
                                                                                                                                    PIDN:AAB89352.1; PID:g264864:
                                                                                                                                                                                                                                                                                                     sulfate-reducing archaeor
<u>..</u>
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Gaps
0
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C;Accession: D96750

C;Accession: D96750

C;Accession: D96750

C;Accession: D96750

R;Theologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White, O.; Alonso, Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; Dewar, K.; Ansen, N.F.; Hughes, B.; Huizar, L.

Nature 408, 816-820, 2000

A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, E.; Kim, C. C.A.; Li, J.H.; Li, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R.; Marziali, Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.; Southwick, A.M.; Sun, H.; Tallon, A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.; Tallon, A;M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.

A.Title: Sequence and analysis of Chromosome 1 of the plant Arabidopsis.

Ference number: A86141; MUID:21016719; PMID:11130712
                                                                                                                                      F;305-321/Domain: transmembrane #status
                                                                                                                                                                                                                                                                                                                                                                                 A;Molecule type: DNA
A;Residues: 1-321 <TIGR>
A;Cross-references: GB:U32787; GB:L42023; NID:g1574619; PIDN:AAC22724.1; PID:g1574620;
A;Note: named as homolog to a protein from Escherichia coli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nrfD protein - Haemophilus influenzae (strain Rd KW20)
C;Species: Haemophilus influenzae
C;Date: 18-Aug-1995 #sequence_revision 26-Jul-1996 #text_change 11-Jun-1999
C;Accession: I64180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 unknown protein F28P22.22 (imported) - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 31-Mar-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
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A; Residues: 1-320 <STO>
                                                                                                                                                                                                                      Description: probably involved in the transfer of electrons from the quinone pool to Superfamily: nrfD protein Reywords: transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nce 269, 496-512, 1995
Thors: Gnehm, C.L.; McDonald, L.A.; Small, K.V.; Fraser, C.M.; Smith, H.O.; Venter, Fritle: Whole-genome random sequencing and assembly of Haemophilus influenzae Rd.; Reference number: A64000; MUID:95350630; PMID:7542800; Accession: I64180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;Fleischmann, R.D.; Adams, M.D.; White, O.; Clayton, R.A.; Kirkness, E.F.; Kerlavage, Gocayne, J.D.; Scott, J.; Shirley, R.; Liu, L.I.; Glodek, A.; Kelley, J.M.; Weidman, D.M.; Brandon, R.C.; Fine, L.D.; Fritchman, J.L.; Fuhrmann, J.L.; Geoghagen, N.S.M.
                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                                                                                    ;Function:
                                                                                                                                                                                                                                                                                                                                              Gene: nrfD
                                                                                                                                                                                                                                                                                                                                                                     Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Status: nucleic acid sequence not shown; translation not shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Genetics
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                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Status: preliminary
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les 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           67 HKKKVAHR
                                                       Local Similarity es 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3 HKSEVAHRFK 12
  1 DAHKSEVAHRFK 12
                                                          Conservative
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                                                                               54.7%;
41.7%;
                                                                                                                                                                    #status predicted #status predicted
                                                       4
                                                                                     Score 35; DB
Pred. No. 85;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 35;
Pred. No.
                                                          Mismatches
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                                                                                                            DB 1; Length 321;
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                                                       Indels
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                                                       Gaps
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R;Taylor, R.G.; Lambert, M.A.; Sexsmith, E.; Sadler, S.J.; Ray, P.N.; Mahuran, D.J.; McIr J. Biol. Chem. 265, 18192-18199, 1990
A;Title: Cloning and expression of rat histidase. Homology to two bacterial histidases ar A;Reference number: A36087; MUID:91009306; PMID:2120224
A;Accession: A36087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Suchi, M.; Harada, N.; Wada, Y.; Takagi, Y.
Biochim. Biophys. Acta 1216, 293-295, 1993
A;Title: Molecular cloning of a cDNA encoding human histidase
A;Reference number: $43415, MUID:94060103, PMID:7916645
A;Accession: $43415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 31
$43415
histidine ammonia-lyase (EC 4.3.1.3) - human
                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: GB:M58308; GB:J05653; NID:g204556; PIDN:AAA63491.1; C;Superfamily: histidine ammonia-lyase C;Keywords: ammonia-lyase; carbon-nitrogen lyase; histidine catabolism F;253-255/Cross-link: 5-imidazolinone (Ala-Gly) #status predicted
                                                                        RESULT 33
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A; Residues: 1-657 <TAY>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Date: 25-Jan-1991 #sequence_revision 25-Jan-1991 #text_change 10-Dec-1999 C;Accession: A36087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C; Species: Rattus norvegicus (Norway rat)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               histidine ammonia-lyase (EC 4.3.1.3) - rat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Superfamily: histidine ammonia-lyase
C;Keywords: ammonia-lyase; carbon-nitrogen lyase; histidine catabolism
F;253-255/Cross-link: 5-imidazolinone (Ala-Gly) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Pathway: histidine catabolism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Description: catalyzes the formation of (E)-3-(1H-imidazol-4-yl)-propenoic acid (urocar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C; Function
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Gene: GDB: HAL; HIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references: DDBJ:D16626; NID:g451209; PIDN:BAA04047.1; PID:g451210
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A; Residues: 1-657 < SUC>
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C;Accession: $43415
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N; Alternate names: histidase
                          histidine ammonia-lyase (EC 4.3.1.3) -
                                                     A46128
                                                                                                                                                                                                                                                                                                                                                                  F;254/Modified site: dehydroalanine (Ser) #status predicted
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                                                                                                                                                                                                                                                                                    Best
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Best Local
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les 8; Conservat
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                                                                                                                                                           DHHPSEIAESHRF 387
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8; Conserv
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                                                                                                                                                                                                                                                         Score 35; DB 2; LC
Pred. No. 1.8e+02;
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Pred. No. 1.8e+02;
                                                                                                                                                                                                                                                                                                               Length 657;
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7:
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C;Species: Mus musculus (house mouse)
C;Date: 21-Sep-1993 #sequence_revision 18-Nov-1994 #text_change 10-Dec-1999
C;Accession: A46128
R;Taylor, R.G.; Grieco, D.; Clarke, G.A.; McInnes, R.R.; Taylor, B.A.
Genomics 16, 231-240, 1993
A;Title: Identification of the mutation in murine histidinemia (his) and genetic mapping A;Reference number: A46128; MUID:93252384; PMID:8486363
R;Watabe, K.; Ishikawa, T.; Mukohara, Y.; Nakamura, H.
J. Bacteriol. 174, 962-969, 1992
A;Title: Cloning and sequencing of the genes involved in the conversion of 5-substituted A;Reference number: A42594; MUID:92121137; PMID:1732229
A;Accession: B42594
                                                                                                              D-amino acid hydantoin hydrolase (ATP-hydrolyzing)
C;Species: Pseudomonas sp.
C;Date: 04-Mar-1993 #sequence_revision 18-Nov-1994
C;Accession: B42594
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Heidelberg, J.F.; Eisen, J.A.; Nelson, W.C.; Clayton, R.A.; Chardson, D.; Ermolaeva, M.D.; Vamathevan, J.; Bass, S.; Qin, I, R.R.; McKalanos, J.J.; Venter, J.C.; Fraser, C.M. Nature 406, 477-483, 2000
                                                                                                                                                                                                                B42594
                                                                                                                                                                                                                                   RESULT 35
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Species: Vibrio cholerae
C;Date: 18-Aug-2000 #sequer
C;Accession: H82143
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-657 <TAY>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      methyl-accepting chemotaxis protein VC1898 [imported] - Vibrio cholerae (strain N16961
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Note: sequence extracted from NCBI backbone (NCBIN:131641, NCBIP:131644)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Date: 18-Aug-2000 #sequence_revision 20-Aug-2000 #text_change 02-Feb-2001
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;Experimental source: C57BL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gene: VC1898
                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Keywords: ammonia-lyase; carbon-nitrogen lyase; histidine catabolism 3-255/Cross-link: 5-imidazolinone (Ala-Gly) #status predicted 4/Modified site: dehydroalanine (Ser) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     oss-references: GB:AE004265; GB:AE003852; NID:g9656424; PIDN:AAF95046.1; GSPDB:GN001
perimental source: serogroup Ol; strain N16961; biotype El Tor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       position: 1
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                                                                                                                                                                                                                                                                                                        663 QSELVHRFK 671
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Pred. No. 1.8e+02;
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Pred. No. 1.8e+02;
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                                                                                                                                   #text_change 16-Feb-2001
                                                                                                                                                                                   (EC 3.5.2.-) hyuA [validated]
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H.; Dragoi, I.; Sell
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A; Map position:
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                                                                                                                                                                                                                                                      A;Gene:
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Best Local
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                          187
                                                                  2 AHKSEVAHRFK 12
                          AHKNSCAHGYK 197
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hypothetical protein K03A11.1 - Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Datc: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 18-Feb-2000
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A;Cross-references: GB:M72717; NID:g151280; PIDN:AAA25845.1; PID:g151282
A;Cross-references: GB:M72717; NID:g151280; PIDN:AAA25845.1; PID:g151282
A;Note: sequence extracted from NCBI backbone (NCBIN:77753, NCBIP:77760)
C;Superfamily: Pseudomonas D-amino acid hydantoin hydrolase (ATP-hydrolyzing) hyuA
C;Keywords: hydrolase
                                                                                                                                                                                                    A;Cross-references: EMBL:Z77133; PIDN:CAB00865.1; GSPDB:GN00028; CESP:K03A11.1
A;Experimental source: clone K03A11
                                                                                                                                                                                                                                                                                                       A; Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                           submitted to the EMBL Data Library, July 1996
A;Reference number: Z19717
A;Accession: T23256
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A; Accession: T38327
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 03-Dec-1999
C;Accession: T38327
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                                                                                                                                                                                                                                                        A;Residues: 1-969 <WIL>
                                                                                                                                                                                                                                                                             A; Molecule type:
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A;Experimental source: strain 972h-; cosmid c23H4
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nes 8; Conserv
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nes 5; Conserv
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                                                                                                                                                      CESP: K03A11.1
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                                                                                                    33/1; 116/1; 162/1; 185/1; 256/2; 359/1; 494/1; 580/1;
                               Similarity
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       Conservative
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45.5%;
                            54.7%; Score 35; DB 2; 54.5%; Pred. No. 2.7e+02;
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Pred. No. 2e+0
0; Mismatches
       2
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Pred. No. 1.9e+02;
3; Mismatches 3
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          Mismatches
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2e+02;
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       Indels
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Gaps
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saframycin Mxl synthetase A - Myxococcus xanthus
(;Species: Myxococcus xanthus
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 17-Nov-2000
C;Accession: T1852
R;Pospiech, A.; Bietenhader, J.; Schupp, T.
Microbiology 142, 741-746, 1996
A;Title: Two multifunctional peptide synthetases and an O-methyltransferase are
A;Reference number: Z18967; MUID:97090395; PMID:8936303
A;Accession: T18552
type II activin receptor ActRIIB - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 19-Dec-1993 #sequence_revision 18-Nov-1994 #text_chan
C;Accession: B49193
R;Feng, Z.M.; Madigan, M.B.; Chen, C.L.
Endocrinology 132, 2593-2600, 1993
A;Title: Expression of type II activin receptor genes in the A;Reference number: A49193; MUID:93279247; PMID:7916681
A;Accession: B49193
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B49193
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Comment: This protein plays a role in lung development and involved in transforming C;Superfamily: activin receptor II; protein kinase homology F;1-251/Domain: protein kinase homology (fragment) <KIN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Zhao, Y.; Silbajoris, R.; Young, S.L.
Biochem. Biophys. Res. Commun. 229, 50-57, 1996
A;Title: Identification and developmental expression of two
A;Reference number: PC4260; MUID:97112402; PMID:8954082
A;Contents: lung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            activin type II receptor - baboon (fragment)
C;Species: Papio sp. (baboon)
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A;Residues: 1-2605 <POS>
A;Cross-references: EMBL:U24657; NID:g1171127; PID:g1171129; PIDN:AAC44129.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Species: Papio sp. (baboon)
;Date: 11-Apr-1997 #sequence_revision 09-May-1997 #text_change 23-Aug-1997
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Best Local :
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Best Local
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97-1065/Domain: acyl carrier protein homology <ACP1>
643-2091/Domain: acetate-CoA ligase homology <ACL2>
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les 5; Conserv
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41.7%;
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Pred. No.
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                                                                                                                                           #text_change 28-Feb-1997
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activin receptor isoform IIB4 - mouse C;Species: Mus musculus (house mouse) C;Date: 04-Mar-1993 #sequence_revision C;Accession: B40829 R;Attisano, L.; Wrana, J.L.; Cheifetz, Cell 68, 97-108, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein disulfide-isomerase (EC 5.3.4.1) - castor bean C;Species: Ricinus communis (Castor bean) C;Date: 19-Mar-1997 #sequence_revision 25-Apr-1997 #text_change C;Accession: S62626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Note: sequence extracted from NCBI backbone (NCBIN:133010, NCBIP:133011)
C;Superfamily: activin receptor II; protein kinase homology
C;Keywords: ATP
F;74-371/Domain: protein kinase homology <KIN>
                                                                                                                                                                                                                                                                        A;Title: Novel activin receptors: distinct genes and alternative A;Reference number: A40829; MUID:92119722; PMID:1310075 A;Accession: B40829
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Molecule type: nucleic acid
A; Residues: 1-382 < FEN>
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                                                                                                        A;Experimental source: Balb/c 3T3
A;Note: sequence inconsistent with the nucleotide translation
A;Note: sequence extracted from NCBI backbone (NCBIN:88364, NN
C;Superfamily: activin receptor II; protein kinase homology
C;Keywords: ATP; receptor
                                                                                                                                                                                                                A; Molecule type: mRNA
A; Residues: 1-504 < ATT>
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A; Residues: 1-498 <COU>
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A;Title: Molecular characterisation of plant endoplasmic A;Reference number: S62620; MUID:96202938; PMID:8631332
A;Accession: S62626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Coughlan, S.J.; Hastings, C.; Winfrey Jr., Eur. J. Biochem. 235, 215-224, 1996
                                                                                                                                                                                                                                                     A; Status: preliminary
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                                                                                    F;180-477/Domain: protein kinase homology <KIN>
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Local Similarity hes 7; Conserv
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      Score 34.5; D
Pred. No. 1.7e
2; Mismatches
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Pred. No. 1.7e+02;
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                            1.7e+02;
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C;Species: Mus musculus (house mouse)
C;Date: 04-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 23-May-1997
C;Accession: D40829
R;Attisano, L.; Wrana, J.L.; Cheifetz, S.; Massague, J.
Cell 68, 97-108, 1992
Cell 68, 97-108 activin receptors: distinct genes and alternative mRNA splicing generate A;Rtele: Novel activin receptors: distinct genes and alternative mRNA splicing generate A;Reference number: A40829; MUID:92119722; PMID:1310075
A;Accession: D40829
A;Status: preliminary
A;Status: preliminary
activin receptor precursor - rat (;Species: Rattus norvegicus (Norway rat))
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 23-May-1997
C;Accession: JQ1484
R;Legerski, R.; Zhou, X.; Dresback, J.; Eberspaecher, H.; McKinney, S.; Segarini, Biochem. Biophys. Res. Commun. 183, 672-679, 1992
A;Title: Molecular cloning and characterization of a novel rat activin receptor. A;Reference number: JQ1484; MUID:92198455; PMID:1312838
A;Accession: JQ1484
A;Molecule type: mRNA
A;Residues: 1-513 <LEG>
C;Superfamily: activin receptor II; protein kinase homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         activin type II receptor - human C;Species: Homo Bapiens (man) C;Decies: Homo Bapiens (man) C;Date: 29-May-1998 #sequence_revision 29-May-1998 #text_change 18-Jun-1999 C;Accession: I37134
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C;Species: Mus musculus (house mouse)
C;Date: 04-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 23-May-1997
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A; Residues: 1-512 < ATT>
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;Residues: 1-512 <RES->
;Residues: 1-512 <RES->
;Cross-references: EMBL:X77533; NID:g825619; PIDN:CAA54671.1;
Cross-references: EMBL:X77533; NID:g825619; PIDN:CAA54671.1;
Poerfamily: activin receptor II; protein kinase homology
6-485/Domain: protein kinase homology <KIN>
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nes 7; Conserv
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Pred. No. 1.7e+02;
2; Mismatches 3;
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A; Status preliminary
A; Status preliminary
A; Molecule type: mRNA
A; Residues: 1-36 < ATT>
A; Cross-references: GB: M84120; NID: g191668; PIDN: AAA37172.1; PID: g191669
A; Cross-references: GB: M84120; NID: g191668; PIDN: AAA37172.1; PID: g191669
A; Note: sequence inconsistent with the nucleotide translation
A; Note: sequence extracted from NCBI backbone (NCBIN: 76259, NCBIP: 76260)
A; Note: sequence extracted from NCBI backbone (NCBIN: 76259, NCBIP: 76260)
C; Superfamily: activin receptor II; protein kinase homology
C; Keywords: ATP; receptor; serine/threonine-specific protein kinase; tran
F; 212-509/Domain: protein kinase homology < KIN>
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A;Note: sequence inconsistent with the nucleotide translation
A;Note: sequence extracted from NCBI backbone (NCBIN:88362, NCBIP:88363)
C;Superfamily: activin receptor II; protein kinase homology
C;Keywords: ATP; receptor
F;204-501/Domain: protein kinase homology <KIN>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Attisano, L.; Wrana, J.L.; Cheifetz, S.; Massague, J. Cell 68, 97-108, 1992
Cell 68, 97-108, 1992
A;Title: Novel activin receptors: distinct genes and alternative mRNA splicing generate & A;Reference number: A40829; MUID:92119722; PMID:1310075
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 activin receptor isoform IIB1 - mouse
C;Species: Mus musculus (house mouse)
C;Date: 04-Max-1993 #sequence_revision 18-Nov-1994 #text_change 18-Jun-1999
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A; Residues: 1-528 < ATT>
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Cell 68, 97-108, 1992
A;Title: Novel activin receptors: distinct genes and alternative mRNA splicing generate and A;Reference number: A40829; MUID:92119722; PMID:1310075
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C;Date: 04-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 23-May-1997
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                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Match 53.9%;
Local Similarity 53.8%;
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   Similarity 7; Conserv
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                            53.9%;
53.8%;
Score 34.5; DB 2;
Pred. No. 1.8e+02;
2; Mismatches 3
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Pred. No. 1.8e+02;
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                                                                DB 2;
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                                                         Length 536;
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1 DAHKSEVAHR-FK 12

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R;Takami, H.; Nakasone, K.; Takaki, Y.; Maeno, G.; Sasaki, R.; Masui, N.; Fuji, F.; Hir: Nucleic Acids Res. 28, 4317-4331, 2000
Nucleic Acids Res. 28, 4317-4331, 2000
A;Title: Complete genome sequence of the alkaliphilic bacterium Bacillus halodurans and A;Reference number: A83650; MUID:20512582; PMID:11058132
A;Accession: C84068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            polyribonucleotide nucleotidyltransferase (general stress protein 13) BH3347 [imported] C;Species: Bacillus halodurans C;Date: 01-Dec-2000 #sequence_revision 01-Dec-2000 #text_change 15-Jun-2001 C;Accession: C84068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 48
C84068
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R;DiSalvo, J.; Bayne, M.L.; Conn, G.; Kwok, P.W.; Trivedi, P.G.; Soderman, D.D.; Palisi J. Biol. Chem. 270, 7717-7723, 1995
A;Title: Purification and characterization of a naturally occurring vascular endothelia
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N;Alternate names: protein F17114.40
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Jun-2000 #sequence_revision 02-Jun-2000 #text_change 02-Sep-2000
                                                                                                                                                                        RESULT 50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     submitted to the Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;Status: preliminary;
Molecule type: DNA
;Residues: 1-156 - SBV>
;Cross-references: EMBL:AL353994; GSPDB:GN00063; ATSP:F17I14.40
perimental source: cultivar Columbia; BAC clone F17I14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bevan, M.; Hilbert, H.; Braun, M.; Holzer, E.; Brandt, A.; Duesterhoeft, A.; Bancroft,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Superfamily: polyribonucleotide nucleotidyltransferase homolog
                                                                                                                                                                                                                                                                                                                                                                                                                     ;Map position: 5
;Introns: 19/2; 37/3; 54/:
;Superfamily: Escherichia
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Accession: T49921
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Residues: 1-138 <STO>
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les 7; Conserv
                                                                                                                                                                                                                                          74 HKIFTELAHRYK 85
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Purification and characterization of a naturally occurring vascular endothelial
                                                                                                                                                                                                                                                                                      HK--SEVAHRFK 12
                                                                                                                                                                                                                                                                                                                            Similarity 58.3%;
7; Conservation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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chia coli ribosomal protein L17
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Pred. No.
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Pred. No. 54;
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A,Status: preliminary; not compared with conceptual translation A,Molecule type: mRNA A,Residues: 1-158 <DIS>
В
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                                                                                                                  C; Keywords: glycoprotein
                                                                                                                                A; Cross-references: GB:L40030; NID:g1263413; PIDN:AAA97426.1; PID:g1263414
                                                                                                                                                                                          A;Reference number: A56125; MUID:95221439; PMID:7706320 A;Accession: A56125
                                                         Best Loc
Matches
                                                                                       Query Match
 59 DEHPNEVSHIF
                                                       Local Similarity
les 6; Conserv
                             1 DAHKSEVAHRF 11
                                                           Conservative
 69
                                                                          53.1%;
54.5%;
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Pred. No.
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62;
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                                                            Gaps
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Search completed: April 11, 2003, 15:19:46
Job time : 18 secs

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 75 summaries
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Perfect score:
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Published Applications AA:*

1: /cgn2 6/ptodata/2/pubpaa/US08 NEW PUB.pep:*

2: /cgn2-6/ptodata/2/pubpaa/PCT NEW PUB.pep:*

3: /cgn2-6/ptodata/2/pubpaa/US06 NEW PUB.pep:*

4: /cgn2-6/ptodata/2/pubpaa/US06 PUBCOMB.pep:*

5: /cgn2-6/ptodata/2/pubpaa/US07 PUBCOMB.pep:*

6: /cgn2-6/ptodata/2/pubpaa/US07 PUBCOMB.pep:*

7: /cgn2-6/ptodata/2/pubpaa/US08 PUBCOMB.pep:*

9: /cgn2-6/ptodata/2/pubpaa/US09 NEW PUB.pep:*

9: /cgn2-6/ptodata/2/pubpaa/US09 NEW PUB.pep:*

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11: /cgn2-6/ptodata/2/pubpaa/US09 PUBCOMB.pep:*

12: /cgn2-6/ptodata/2/pubpaa/US09 PUBCOMB.pep:*

13: /cgn2-6/ptodata/2/pubpaa/US09 PUBCOMB.pep:*

14: /cgn2-6/ptodata/2/pubpaa/US09 PUBCOMB.pep:*

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16: /cgn2-6/ptodata/2/pubpaa/US09 PUBCOMB.pep:*

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64
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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651	610	610	610	610	610	609	585	585	268	241	195	26	24	17	13	12	12	12	Match Length
7.	10	g	9	9	9	12	12	10	9	9	9	9	9	9	9	9	9	9	8
US-10-153-064-133	US-09-984-186-2	US-10-237-871-2	US-10-237-866-2	US-10-237-708-2	US-10-237-667-2	US-10-153-064-7	US-10-153-064-5	US-09-929-552-2	US-10-074-956-28	US-10-074-956-27	US-10-074-956-24	US-09-846-329-1	US-09-846-328-1	US-09-845-727-1	US-09-845-764-1	US-10-076-071-6	US-10-076-071-3	US-09-846-347-1	ID
sequence 133, App	Sequence 2, Appli	Sequence 7, Appli	Sequence 5, Appli	Sequence 2, Appli	Sequence 28, Appl	•	Sequence 24, Appl	Sequence 1, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 6, Appli	Sequence 3, Appli	Sequence 1, Appli	Description				

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Sequence 4, Appli Sequence 36, Appl Sequence 37, Appl Sequence 30, Appl Sequence 32, Appl Sequence 34, Appl Sequence 37, Appl Sequence 37, Appl Sequence 39, Appli Sequence 39, Appli	C C C C C C C C C C C C C C C C C C C	Sequence 96, Appl

ALIGNMENTS

RESULT 1 US-09-846-347-1

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Sequence 1, Application US/09846347;
Publication No. US20030040602A1
GENERAL INFORMATION:
APPLICANT: Jackowski, George
TITLE OF INVENTION: BIOPOLYMER MARKER INDICATIVE OF DISEASE STATE HAVING A MOLECULAR TITLE OF INVENTION: OF 1406 DALTONS
FILE REFERENCE: 2132.032
CURRENT APPLICATION NUMBER: US/09/846,347;
CURRENT FILING DATE: 2001-04-30
NUMBER OF SEQ ID NOS: 1
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Sequence
Publication No. ...
GENERAL INFORMATION:
GAPPLICANT: Bar-Or, David
Curtis, C. Ger
Tan, Edward
Tan, Edward
                                                                                                                                                                            US-10-076-071-6
                                                                                                                                                                                                RESULT 3
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                                                                                                                                                                                                                                                                                                                                                                                                        GRGANISM: Homo sapiens US-10-076-071-3
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PRIOR APPLICATION NUMBER: 09/678,202
PRIOR FILING DATE: 2000-09-29
PRIOR APPLICATION NUMBER: 60/283,507
PRIOR APPLICATION NUMBER: 09/281,679
PRIOR PILING DATE: 2001-03-22
PRIOR FILING DATE: 2001-03-22
                                                                                                                                     Sequence 6, Application US/10076071 Publication No. US20030060408A1
                                                                                                                                                                                                                                                                                                                          Query Match 100.0%; Score 64; DB 9; 18est Local Similarity 100.0%; Pred. No. 6.8e-06; Matches 12; Conservative 0; Mismatches 0;
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Publication No-US/20030060408A1
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LENGTH: 12
TYPE: PRT
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APPLICANT: Curti
APPLICANT: Lau,
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PRIOR FILING DATE: 1999-10-01
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TITLE OF INVENTION: Metal-Binding Compounds and Uses Therefor
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Lau, Edward
Rao, Nagaraja K.R.
Winkler, James V.
Crook, Wannell M.
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Lau, Edward
Rao, Nagaraja K.R.
Winkler, James V.
Crook, Wannell M.
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Pred. No. 6.8e-06;
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RESULT 5
US-09-845-727-1
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APPLICANT: Jackowski, George
TITLE OF INVENTION: BIOPOLYMER MARKER INDICATIVE OF DISEASE STATE HAVING A MOLECULAR V
TITLE OF INVENTION: OF 1521 DALTONS
                                                                                                                                                          Query Match
Best Local Similarity
Matches 12; Conserv
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NUMBER OF SEQ ID NOS: 9
SOFTWARE: PatentIn version 3.0
SEQ ID NO 6
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CURRENT FILING DATE: 2001-04-30
NUMBER OF SEQ ID NOS: 1
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PRIOR FILING DATE: 1999-10-01
PRIOR APPLICATION NUMBER: 60/157,404
PRIOR FILING DATE: 1999-10-01
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PRIOR FILING DATE: 2000-09-29
PRIOR APPLICATION NUMBER: 60/283,507
PRIOR FILING DATE: 2001-04-11
PRIOR APPLICATION NUMBER: 09/816,679
PRIOR FILING DATE: 2001-03-22
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CURRENT APPLICATION NUMBER: US/10/076,071
CURRENT FILING DATE: 2002-02-13
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                                                                                                                                                                                                                                                                                         LENGTH: 13
TYPE: PRT
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ORGANISM: Homo mapiens
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NAME/KEY: MOD RES
LOCATION: (1)...(1
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les 12; Conserv
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100.0%; Pred. No. 6.8e-06;
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Sequence 1, Application US/09845727 Patent No. US20020160418A1 GENERAL INFORMATION:

APPLICANT: Jackowski, George

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; SEQ ID NO 1; LENGTH: 24; TYPE: PRT; ORGANISM: Homo sapiens US-09-846-328-1
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US-09-846-328-1
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Patent No. US20020160531A1
GENERAL INFORMATION:
APPLICANT: Jackowski, George
APPLICANT: Jackowski, George
TITLE OF INVENTION: BIOPOLYMER MARKER INDICATIVE OF DISEASE STATE HAVING A MOLECULAR
TITLE OF INVENTION: OF 2753 DALTONS
FILE REFERENCE: 2132.051
CURRENT APPLICATION NUMBER: US/09/846,328
CURRENT FILING DATE: 2001-04-30
                                                                                                                                                                                NUMBER OF SEQ ID NOS: 1
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1
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Patent No. US20020161177A1
GENERAL INFORMATION:
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                                                              Query Match
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TITLE OF INVENTION: Blopolymer Marker Indicative of Disease State Having A Molecular
TITLE OF INVENTION: of 2937 Daltons
FILE REFERENCE: 2132.052
CURRENT APPLICATION NUMBER: US/09/846,329
CURRENT FILING DATE: 2001-04-30
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SOFTWARE: PatentIn version 3.1
                                                                                                                     LENGTH: 26
TYPE: PRT
ORGANISM: Homo sapiens
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                    y Match 100.0%; Score 64; DB 9; 1
Local Similarity 100.0%; Pred. No. 1.6e-05;
hes 12; Conservative 0; Mismatches 0;
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                                                            DB 9; Length 26;
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; LENGTH: 195
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-074-956-24
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Sequence 28, Application US/10074956
Publication No. US2002019332A1
GENERAL INFORMATION:
APPLICANT: Hedley, Mary Lynne
TITLE OF INVENTION: METHODS OF TREATING BLADDER DISORDERS
FILE REFERENCE: 08191-022001
CURRENT APPLICATION UMBER: US/10/074,956
CURRENT FILING DATE: 2002-06-10
                                                                                                                                                                                                                                                                                                                                                                                                       ; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-074-956-27
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US-10-074-956-24
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Best Local S
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Sequence 27, Application US/10074956 Publication No. US20020193332A1 GENERAL INFORMATION:
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TITLE OF INVENTION: METHODS OF TREATING BLADDER DISORDERS
FILE REFERENCE: 06191-022001
CURRENT APPLICATION NUMBER: US/10/074,956
CURRENT FILING DATE: 2002-06-10
PRIOR APPLICATION NUMBER: 60/268,175
PRIOR APPLICATION UMBER: 60/268,175
PRIOR FILING DATE: 2001-02-12
NUMBER OF SEQ ID NOS: 29
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PRIOR FILING DATE: 2001-02-12
NUMBER OF SEQ ID NOS: 29
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CURRENT APPLICATION NUMBER: US/10/074,956
CURRENT FILING DATE: 2002-06-10
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y Match 100.0%;
Local Similarity 100.0%;
hes 12; Conservative 0
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Score 64; DB 9;
Pred. No. 0.0002;
); Mismatches 0
                                                      Length 241;
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-074-956-28
                                                                                                                                                  US-09-929-552-2
                                                                          Matches
                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Patent No. US20020123080A1
GENERAL INFORMATION:
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Best Local Similarity
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PRIOR FILING DATE: 2001-02-12
NUMBER OF SEQ ID NOS: 29
                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 2:
                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25 DAHKSEVAHRFK 36
                       1 DAHKSEVAHRFK 12
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
                                                                                                                                                             TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/929,552
FILING DATE: 14-Aug-2001
CLASSIFICATION: cUnknowns
PRIOR APPLICATION DATA: 08/769,746
APPLICATION NUMBER: 08/769,746
ATIONG DATE: 19-DEC-1996
ATTORNETY/AGENT INFORMATION:
                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 585 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Medlen & Carroll, LLP
                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: MBRI-02584 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Soto, Ana M.
TITLE OF INVENTION: Inhibiting Proliferation of Cancer Cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Sonnenschein, Carlos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
DAHKSEVAHREK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12;
                                                                                                                                                                                                                                                                                                                                                                        NAME: Carroll, Peter G. REGISTRATION NUMBER: 32,837
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                              TELEFAX: (415) 397-8338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: United States of America
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                                                                        Conservative
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                                                                                     100.0%;
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                                                                      Score 64; DB 10;
Pred. No. 0.00053;
; Mismatches 0;
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Pred. No. 0.00022;
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                                                                                                         DB 10; Length 585;
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RESULT 12

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RESULT 14
US-10-237-667-2
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US-10-153-064-7
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APPLICANT: Bell et al.
TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: PF556
CURRENT APPLICATION NUMBER: US/10/153,064
CURRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR FILING DATE: 2001-05-25
PRIOR FILING DATE: 2001-05-25
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Patent No. US20020142814A1
GENERAL INFORMATION:
APPLICANT: Bell et al.
TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: PF556
                                                                                                                                                                                                 Sequence 2, Application US/10237667
Publication No. US20030022308A1
GENERAL INFORMATION:
APPLICANT: Fleer, Reinhard
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SEQ ID NO 5
LENCTH: 585
TYPE: PRT
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Patent No. US20020142814A1
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PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
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CURRENT FILING DATE: 2002-05-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Homo Sapiens
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les 12; Conservative
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NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Rhone-Poulenc Rorer Inc
                                                                                                     TITLE OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DAHKSEVAHRFK 12
                                   Jung, Gerard
Yeh, Patrice
Yeh, Patrice
INVENTION: NOTE BIOLOGICALLY ACTIVE POLYPEPTIDES,
PREPARATION THEREOF AND PHARMACEUTICAL COMPOSITION
PROPERTY SAID POLYPEPTIDES
                                                                                                                                                           Fournier, Alain
Guitton, Jean-Dominique
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Pred. No. 0.00053;
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500 Arcola Road, 3C43

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LENGTH: 610 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 2:
US-10-237-667-2
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Best Local Similarity 100.0%; Pred. No. 0.00055;
Matches 12; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/10237708 Publication No. US20030036170A1 GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/797,689
FILING DATE: 31-JAN-1997
APPLICATION NUMBER: US 08/256,927
FILING DATE: 28-JUL-1994
APPLICATION NUMBER: FR 92/01064
APPLICATION NUMBER: FR 92/01064
APILING DATE: 31-JAN-1992
APPLICATION NUMBER: PCT/FR93/00085
FILING DATE: 28-JAN-1993
ATTORNEY/AGENT INFORMATION:
NAME: Smith Ph.D., Julie K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25 DAHKSEVAHRFK 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 DAHKSEVAHRFK 12
COUNTRY: USA
ZIP: 19426
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
OPERATING SYSTEM: System 7.1
                                                                                                                                                                                                                                                                               APPLICANT: Fleer, Reinhard
Fournier, Alain
Guitton, Jean-Dominique
Jung, Gerard
Yeh, Patrice
TITLE OF INVENTION: NOVEL BIOLOGICALLY ACTIVE POLYPEPTIDES,
PREPARATION THEREOF AND PHARMACEUTICAL COMPOSITION
                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Rhone-Poulenc Rorer Inc.
STREET: 500 Arcola Road, 3C43
CITY: Collegeville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: P-38,619
REFERENCE/DOCKET NUMBER: ST92006-US
TELECOMMUNICATION INFORMATION:
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FILING DATE: 10-Sep-2002
CLASSIFICATION: cUnknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 19426
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
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OPERATING SYSTEM: System 7.1
SOFTWARE: Word 5.1 (PatentIn
                                                                                                                                      STATE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 500 Arcola CITY: Collegeville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE:
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                                                                                                                                                                                                                                                                      CONTAINING SAID POLYPEPTIDES
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US-10-237-866-2
Sequence 2, Application US/10237866
Publication No. US20030036171A1
GENERAL INFORMATION:
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
COUNTRY: USA
ZIP: 19426
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
OPERATING SYSTEM: System 7.1
SOFTWARE: Word 5.1 (Patentin)
CURRENT APPLICATION NUMBER: US/10/237,866
FILING DATE: 10-5ep-2002
CLASSIFICATION: <Urknown>
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/797,689
FILING DATE: 31-JAN-1997
APPLICATION NUMBER: US 08/256,927
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FILING DATE: 31.JAN-1997
APPLICATION NUMBER: US 08/256,927
FILING DATE: 28.JUL-1994
APPLICATION NUMBER: FF 92/01064
FILING DATE: 31.JAN-1992
APPLICATION NUMBER: PCT/FF93/00085
FILING DATE: 28.JAN-1993
ATTORNEY/AGENT INFORMATION:
NAME: Smith Ph.D., Julie K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25 DAHKSEVAHRFK 36
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                                                                                                                                                                                                                                                                                                                                                                                                              Jung, Gerard
Yeh, Patrice
TITLE OF INVENTION: NOVEL BIOLOGICALLY ACTIVE POLYPEPTIDES,
PREPARATION THEREOF AND PHARMACEUTICAL COMPOSITION
CONTAINING SAID POLYPEPTIDES
                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSE: Rhome-Poulenc Rorer Inc.
STREET: 500 Arcola Road, 3C43
CITY: Collegeville
STATE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Fleer, Reinhard
Fournier, Alain
Guitton, Jean-Dominique
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: P-38,619
REFERENCE/DOCKET NUMBER: ST92006-US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: <Unknown> PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Word 5.1 (PatentIn) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/10/237,708 FILING DATE: 10-Sep-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
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TELEFAX: (610) 454-3808
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100.0%; Pred. No. 0.00055;
tive 0; Mismatches 0;
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FILING DATE: 28-JUL-1994 APPLICATION NUMBER: FR 92/01064

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Publication No. US20030036172A1
GENERAL INFORMATION:
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                                                     PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/797,689
FILING DATE: 31-DAN-1997
APPLICATION NUMBER: US 08/256,927
FILING DATE: 28-UUL-1994
APPLICATION NUMBER: FR 92/01064
FILING DATE: 31-DAN-1992
APPLICATION NUMBER: PCT/FR93/00085
FILING DATE: 28-DAN-1993
ATTORNEY/AGENT INFORMATION:
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25 DAHKSEVAHRFK 36
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mes 12; Conserv
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Rhone-Poulenc Rorer Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 31-JAN-1992
APPLICATION NUMBER: PCT/FR93/00085
FILING DATE: 28-JAN-1993
ATTORNEY/AGENT INFORMATION:
NAME: Smith Ph.D., Julie K.
                                                                                                                                                                                                                                                           COMPUTER: Macintosh
OPERATING SYSTEM: System 7.1
OPERATING SYSTEM: (Patcentin)
CORRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/237,871
PILING DATE: 10-Sep-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jung, Gerard
Yeh, Patrice
TITLE OF INVENTION: NOVEL BIOLOGICALLY ACTIVE POLYPEPTIDES,
PREPARATION THEREOF AND PHARMACEUTICAL COMPOSITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: P-38,619
REFERENCE/DOCKET NUMBER: ST92006-US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
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TELEFAX: (610) 454-3808
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 500 Arcola CITY: Collegeville
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                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
REGISTRATION NUMBER: P-38,619
REFERENCE/DOCKET NUMBER: ST92006-US
                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 19426
                                       NAME: Smith Ph.D., Julie K.
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ilarity 100.0%;
Conservative 0
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Guitton, Jean-Dominique
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US-09-984-186-2
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Best Local Similarity
Matches 12; Conserv
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                 INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                           PRIOR APPLICATION DATA:

APPLICATION UNMEER: US/08/797,689
FILING DATE: 31-JAN-1997
APPLICATION NUMBER: US 08/256,927
FILING DATE: 28-JUL-1994
APPLICATION NUMBER: FR 92/01064
FILING DATE: 31-JAN-1992
APPLICATION NUMBER: PCT/FR93/0085
FILING DATE: 28-JAN-1993
ATTORNAEY/AGENT INFORMATION:
NAME: Smith Ph. D. Julie K.
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Yeh, Patrice
TITLE OF INVENTION: NOVEL BIOLOGICALLY ACTIVE POLYPEPTIDES,
PREPARATION THEREOF AND PHARMACEUTICAL COMPOSITION
CONTAINING SAID POLYPEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: Macintosh
OPERATING SYSTEM: System 7.1
OPERATING SYSTEM: System 7.1
SOFTWARE: WOR'D 5.1 (Patentin)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/984,186
FILING DATE: 29-Oct-2001
CLASSIFICATION: Unknown>
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TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Rhone-Poulenc Rorer Inc.
STREET: 500 Arcola Road, 3C43
CITY: Collegeville
STATE: PA
                                                                                                                                                               REGISTRATION NUMBER: P-38,619
REFERENCE/DOCKET NUMBER: ST92006-US
TELECOMMUNICATION INFORMATION:
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MOLECULE TYPE: protein
                 LENGTH: 610 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 19426
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                                                                                                                           TELEPHONE: (610) 454-3839
TELEFAX: (610) 454-3808
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Guitton, Jean-Dominique
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Pred. No. 0.00055;
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Sequence 96, Application US/10153064

PAtent No. US20020142814A1

GENERAL INFORMATION:

APPLICANT: Bell et al.

TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: FP556

CURRENT APPLICATION NUMBER: US/10/153,064

CURRENT FILING DATE: 2002-05-24

PRIOR APPLICATION NUMBER: 60/293,212

PRIOR FILING DATE: 2001-05-25

NUMBER OF SEQ ID NOS: 137

SOFTMARE: Patentin version 3.1

TYPE: PRT

TYPE: PRT

TYPE: PRT

TYPE: PRT

TYPE: PRT
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; Patent No. US20020142814A1
; GENERAL INFORMATION:
; APPLICANT: Bell et al.
; TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
ILLE REFERENCE: PF556
URRENT APPLICATION NUMBER: US/10/153,064
; CURRENT FILING DATE: 2002-05-24
; PRIOR APPLICATION NUMBER: 60/293,212
; PRIOR FILING DATE: 2001-05-25
; NUMBER OF SEQ ID NOS: 137
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 133
; LENGTH: 651
               RESULT 21
US-10-153-064-99
 ; Sequence
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                                                                                                                                                                                                                                           ; ORGANISM: Homo sapiens
US-10-153-064-96
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US-10-153-064-96
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                                                                                                                                                                    Matches
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Best Local Similarity 100.0%;
Matches 12; Conservative 0
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                                                                                                                                                                                 Local Similarity
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                                                                                            DAHKSEVAHRFK 78
99, Application US/10153064
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                                                                                                                                                                    Conservative
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                                                                                                                                                              100.0%; Score 64; DB 12; Length 652; 100.0%; Pred. No. 0.0006; tive 0; Mismatches 0; Indels
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Pred. No. 0.00055;
; Mismatches 0;
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GENERAL INCOMPATION:
GENERAL INCOMPATION:
APPLICANT: Bell et al.
TITLE OF INVENTION: Chemokine Beta-1 Fusion:
FILE REFERENCE: PF556
CURRENT APPLICATION NUMBER: US/10/153,064
CURRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR FILING DATE: 2001-05-25
INUMBER OF SEQ ID NOS: 137
SOFTWARE: Patentin version 3:1
SEQ ID NO 105
LENGTH: 652
TYPE: PRT
ORGANISM: Homo sapiens
US-10-153-064-105
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APPLICANT: Bell et al.

TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins

FILE REFERENCE: PF556

CURRENT APPLICATION NUMBER: US/10/153,064

CURRENT FILING DATE: 2002-05-24

PRIOR APPLICATION NUMBER: 60/293,212

PRIOR APPLICATION NUMBER: 60/293,212

PRIOR FILING DATE: 2001-05-25

NUMBER OF SEQ ID NOS: 137

SOFTWARE: Patentin version 3.1
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US-10-153-064-105
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                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                           CURRENT APPLICATION NUMBER: US/10/153,064
CURRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
SOFTWARE: Patentin version 3.1
SEQ ID NO 132
SEQ ID NO 132
                                                                                                                                                                                                                                                                                                           Sequence 132, Application US/10153064 Patent No. US20020142814A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local Similarity 100.0%;
Matches 12; Conservative 0
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                                                                                                                                                                                                                    APPLICANT: Bell et al.
TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: PF556
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                    LENGTH: 652
TYPE: PRT
ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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Pred. No. 0.0006;
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CURRENT APPLICATION NUMBER: US/10/153,064
CURRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR FILING DATE: 2001-05-25
INUMBER OF SEO ID NOS: 137
SOFTWARE: Patentin version 3.1
SEO ID NO 131
I SEO ID NO 131
I LENGTH: 653
TYPE: PRT
ORGANISM: Homo sapiens
US-10-153-064-131
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US-10-153-064-130
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                               US-10-153-064-90
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Best Local Similarity 100.
Matches 12; Conservative
                                               RESULT 26
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Patent No. US20020142814A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                       APPLICANT: Bell et al.
TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: PF556
URRENT APPLICATION NUMBER: US/10/153,064
URRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR PRILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
SOFTWARE: Patentin version 3.1
SEQ ID NO 130
Sequence 90, Application US/10153064
Patent No. US20020142814A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                     PATENT NO. US20020142814A1
GENERAL INFORMATION:
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TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: PF556
                                                                                                                                                                              Best
                                                                                                                                                                                                                                           LENGTH: 656
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            68 DAHKSEVAHREK 79
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                                                                                                                               1 DAHKSEVAHRFK 12
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Pred. No. 0.0006;
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PRIOR APPLICATION NUMBER: 60/293,212
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
SOFTWARE: PatentIn version 3.1
SEQ ID NO 90
LENGTH: 660
TYPE: PRT
ORGANISM: Homo sapiens
US-10-153-064-90
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US-10-153-064-93
; Sequence 93, Application US/10153064
; Patent No. US20020142814A1
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-153-064-93
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US-10-153-064-102
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APPLICANT: Bell et al.
TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: PF556
CURRENT APPLICATION NUMBER: US/10/153,064
CURRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
SOFTWARE: Patentin version 3.1
SEQ ID NO 93
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Best Local Similarity
"htches 12; Conserv
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                                                    SOFTWARE: PatentIn version 3.1
SEQ ID NO 102
LENGTH: 668
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TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: PF556
CURRENT APPLICATION NUMBER: US/10/153,064
CURRENT FILING DATE: 2002-05-24
                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                             APPLICANT: Bell et al.
TITLE OF INVENTION: Chemokine Beta-1 Fusion
FILE REFERENCE: PF556
CURRENT APPLICATION NUMBER: US/10/153,064
CURRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
ORGANISM: Homo sapiens FEATURE:
                                    TYPE: PRT
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Pred. No. 0.0006;
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GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
CHECK COP INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: PF556
CURRENT APPLICATION NUMBER: US/10/153,064
CURRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR PILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
SOFTWARE: Patentin version 3.1
SEQ ID NO 95
LENGTH: 676
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                            ; TYPE: PRT ; ORGANISM: Homo sapiens US-10-153-064-98
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                                                             Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                SOFTWARE: PatentIn version 3.1
SEQ ID NO 98
LENGTH: 676
                                                                                                                                                                                                                                      APPLICANT: Bell et al.

APPLICANT: Bell et al.

TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: PF556
CURRENT APPLICATION NUMBER: US/10/153,064
CURRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
NUMBER OF SEQ ID NOS: 137
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Patent No. US20020142814A1
RENERAL INFORMATION:
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91 DAHKSEVAHREK 102
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                                                                 100.0%; Score 64; DB 12; llarity 100.0%; Pred. No. 0.00062; Conservative 0; Mismatches 0;
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                                                                                                     DB 12; Length 676;
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RESULT 31

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Sequence 127, Application US/10153064

Patent No. US20020142814A1

GENERAL INFORMATION:
APPLICANT: Bell et al.

TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: PF556

FILE REFERENCE: P556

CURRENT APPLICATION NUMBER: US/10/153,064

CURRENT FILING DATE: 2002-05-24

PRIOR APPLICATION NUMBER: 60/293,212

PRIOR APPLICATION NUMBER: 60/293,212

PRIOR FILING DATE: 2001-05-25

INUMBER OF SEQ ID NOS: 137

COCUMBER OF SEQ ID NOS: 137
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TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: PF556

CURRENT APPLICATION NUMBER: US/10/153,064

CURRENT FILING DATE: 2002-05-24

PRIOR APPLICATION NUMBER: 60/293,212

PRIOR FILING DATE: 2001-05-25

NUMBER OF SEQ ID NOS: 137

SOFTWARE: Patentin version 3.1

SEQ ID NO 104

LENGTH: 676
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CURRENT APPLICATION NUMBER: US/10/153,064
CURRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
SOFTWARE: Patentin version 3.1
SEQ ID NO 129
LENGTH: 676
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GENERAL INFORMATION:
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Best Local Similarity 100.0%; Pred. No.
Matches 12; Conservative 0; Mismatch
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Patent No. US20020142814A1
                                                                                                                                                                                             APPLICANT: Bell et al.
TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: PF556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                             92 DAHKSEVAHRFK 103
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PRIOR APPLICATION UNMBER 60/293,212
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
SOFTWARE: PattentIn version 3.1
SEQ ID NO 123
LENGTH: 680
TYPE: PRT
ORGANISM: Homo sapiens
US-10-153-064-123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Bell et al.
TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: PF556
CURRENT APPLICATION NUMBER: US/10/153,064
CURRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
SOFTWARE: PatentIn version 3.1
SEQ ID NO 125
LENGTH: 677
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TYPE: PRT ; ORGANISM: Homo sapiens US-10-153-064-129
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 uvery match
Best Local Similarity 100.0%;
Matches 12; Conservative 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 125, Application US/10153064 Patent No. US20020142814A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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                                                                    Matches
                                                                                     Query Match
Best Local S
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                                                                                                                                                                                                                                                                                                    APPLICANT: Bell et al.
ITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
ILE REFERENCE: PF556
CURRENT APPLICATION NUMBER: US/10/153,064
CURRENT FILING DATE: 2002-05-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               watch 100.0%; Score 64; DB 12; Local Similarity 100.0%; Pred. No. 0.00062; nes 12; Conservative 0; Mismatches 0;
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96 DAHKSEVAHRFK 107
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                                   1 DAHKSEVAHRFK 12
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                                                                    12;
                                                                        h 100.0%; Score 64; DB 12; Similarity 100.0%; Pred. No. 0.00062; 0; Mismatches 0;
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Pred. No. 0.00062;
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                                                                                                          DB 12; Length 680;
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RESULT 36 US-10-153-064-92

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US-10-237-667-16
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Patent No. US20020142814A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 92, Application US/10153064
Patent No. US20020142814A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.1 SEQ ID NO 101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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APPLICANT: Bell et al.
TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: PF556
CURRENT APPLICATION NUMBER: US/10/153,064
CURRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                              Sequence 16, Application US/10237667 Publication No. US20030022308A1
                                                                                                                                                                                                                                                                                                                                                                                   Matches
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Best Local
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CURRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
NAME/KEY: SITE
LOCATION: (585)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Bell et al.
TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: PF556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 684
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 692
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                      107 DAHKSEVAHREK 118
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                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
es 12; Conserv
TITLE OF INVENTION: NOVEL BIOLOGICALLY ACTIVE POLYPEPTIDES,
PREPARATION THEREOF AND PHARMACEUTICAL COMPOSITION
CONTAINING SAID POLYPEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12;
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                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                        Jung, Gerard
Yeh, Patrice
                                                                                              Fournier, Alain
Guitton, Jean-Dominique
                                                                                                                                        Fleer, Reinhard
                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
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RESULT 39
US-10-237-708-16
; Sequence 16, Application US/10237708
; Publication No. US20030036170A1
; GENERAL INFORMATION:
APPLICANT: Fleer, Reinhard
. APPLICANT: Fournier, Alain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Thes 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/797,689
PILLING DATE: 31-JAN-1997
APPLICATION NUMBER: US 08/256,927
FILLING DATE: 28-JUL-1994
APPLICATION NUMBER: FR 92/01064
FILLING DATE: 31-JAN-1992
APPLICATION NUMBER: PCT/FR93/00085
FILLING DATE: 28-JAN-1993
ATTORNEY,AGENT INFORMATION:
NAME: Smith Ph.D., Julie K.
REGISTRATION NUMBER: P-38,619
REFERENCE/DOCKET NUMBER: ST92006-US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          203 DAHKSEVAHREK 214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 DAHKSEVAHRFK 12
CITY: Collegeville
STATE: PA
COUNTRY: USA
ZIP: 19416
COMPUTER READABLE FORM:
                                                                                                                         Jung, Gerard
Yeh, Patrice
Yeh, Patrice Novel BIOLOGICALLY ACTIVE POLYPEPTIDES,
PREPARATION THEREOF AND PHARMACEUTICAL COMPOSITION
                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Rhone-Poulenc Rorer Inc.
STREET: 500 Arcola Road, 3C43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 16:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PILING DATE: 10-Sep-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Word 5.1 (Patentin)
CURRENT APPLICATION NUMBER: US/10/237,667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 19426
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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ADDRESSEE: Rhone-Poulenc Rorer Inc.
STREET: 500 Arcola Road, 3C43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 787 amino acids TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (610)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (610) 454-3839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                             Fournier, Alain
Guitton, Jean-Dominique
Jung, Gerard
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Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               203 DAHKSEVAHREK 214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 DAHKSEVAHRFK 12
 PRIOR APPLICATION DATA:
                                                                                                                                                                                                      CITY: Collegeville
STATE: PA
COUNTRY: USA
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Sequence 16, Application US/10237866 Publication No. US20030036171A1 GENERAL INFORMATION:
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TELEPAX: (610) 454-3
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
ZIP: 194.6

ZIP: 194.6

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
OPERATING SYSTEM: System 7.1
SOFTWARE: Word 5.1 (PatentIn)
CURRENT APPLICATION NOMBER: US/10/237,866
FILING DATE: 10-Sep-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/797,689
FILING DATE: 31-JAN-1997
APPLICATION NUMBER: US 08/256,927
FILING DATE: 28-JUL-1994
APPLICATION NUMBER: FR 92/01064
FILING DATE: 31-JAN-1992
APPLICATION NUMBER: PCT/FR93/00085
FILING DATE: 28-JAN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
CORDESSEE Rhome-Poulenc Rorer Inc.
STREET: 500 Arcola Road, 3C43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 16:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: P-38,619
REFERENCE/DOCKET NUMBER: ST92006-US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 10-Sep-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: NOVEL BIOLOGICALLY ACTIVE POLYPEPTIDES,
PREPARATION THEREOF AND PHARMACEUTICAL COMPOSITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/237,708
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Fleer, Reinhard
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Smith Ph.D., Julie K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 787 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fournier, Alain
Guitton, Jean-Dominique
Jung, Gerard
Yeh, Patrice
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y 100.0%; Pr
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Pred. No. 0.0
0; Mismatches
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Sequence 16, Application US/10237871 Publication No. US20030036172A1 GENERAL INFORMATION:
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MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 16:
                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
OPERATING SYSTEM: System 7.1
SOFTWARE: Word 5.1 (Patentin)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Rhone-Poulenc Rorer Inc
STREET: 500 Arcola Road, 3C43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: NOVEL BIOLOGICALLY ACTIVE POLYPEPTIDES,
PREPARATION THEREOF AND PHARMACEUTICAL COMPOSITION
CONTAINING SAID POLYPEPTIDES
    APPLICATION NUMBER: US/08/797,689
FILING DATE: 31-JAN-1997
APPLICATION NUMBER: US 08/256,927
FILING DATE: 28-JUL-1994
APPLICATION NUMBER: FR 92/01064
FILING DATE: 31-JAN-1992
APPLICATION NUMBER: PCT/FR93/00085
FILING DATE: 28-JAN-1993
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FILING DATE: 31-JAN-1997
APPLICATION NUMBER: US 08/256,927
FILING DATE: 28-JUL-1994
                                                                                                                                                                                      APPLICATION NUMBER: US/10/237,871 FILING DATE: 10-Sep-2002 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fournier, Alain
Guitton, Jean-Dominique
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Pred. No. 0.00074;
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US-09-984-186-16
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Best Local Similarity
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            TELEPHONE: (610) 454-38
TELEFAX: (610) 454-3808
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                    APPLICATION NUMBER: US 08/256,927
FILING DATE: 28-JUL-1994
APPLICATION NUMBER: FR 92/01064
FILING DATE: 31-JAN-1992
APPLICATION NUMBER: PCT/FR92/00085
FILING DATE: 28-JAN-1993
ATTORNEY/AGENT INFORMATION:
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REGISTRATION NUMBER: P-38,619
REFERENCE/DOCKET NUMBER: ST92006-US
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: Macintosh
OPERATING SYSTEM: System 7.1
SOFTWARE: WORD 5.1 (Patentin)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/984,186
FILING DATE: 29-Oct-2001
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Rhone-Poulenc Rorer Inc.
STREET: 500 Arcola Road, 3C43
CITY: Collegeville
                                                                      NAME: Smith Ph.D., Julie K.
REGISTRATION NUMBER: P-38,619
REFERENCE/DOCKET NUMBER: ST92006-US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (610) 454-3839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: NOVEL BIOLOGICALLY ACTIVE POLYPEPTIDES,
PREPARATION THEREOF AND PHARMACEUTICAL COMPOSITION
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TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/797,689 FILING DATE: 31-JAN-1997
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LENGTH: 787 amino acids
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Guitton, Jean-Dominique
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100.0%; Pred. No. 0.00074;
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Best Matches

Local Similarity

Conservative

100.0%; 0

Query Match

US-10-237-866-16

LENGTH: 787 amino acids TYPE: amino acid

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203 DAHKSEVAHREK 214

1 DAHKSEVAHREK 12

RESULT 41 US-10-237-871-16

APPLICANT: Fleer, Reinhard

Jung, Gurice

NUMBER OF SEQUENCES:

CITY: Collegeville STATE: PA

ZIP: COUNTRY:

19426

USA

PRIOR APPLICATION DATA:

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US-10-153-064-89

** Sequence 89, Application US/10153064

**Patent No. US20020142814A1

** GENERAL INFORMATION:

** APPLICANT: Bell et al.

** TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins

** FILE REFERENCE: PF556

** CURRENT APPLICATION NUMBER: US/10/153,064

** CURRENT FILLING DATE: 2002-05-24

** PRIOR APPLICATION NUMBER: 60/293,212

** PRIOR PILING DATE: 2001-05-25
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PRIOR FILING DATE: 1998-01-08
PRIOR FILING DATE: 1998-06-07
PRIOR FILING DATE: 1995-06-07
PRIOR APPLICATION NUMBER: 08/295,078
PRIOR APPLICATION NUMBER: 08/295,078
PRIOR FILING DATE: 1994-08-26
PRIOR PRIOR APPLICATION NUMBER: 08/121,236
PRIOR FILING DATE: 1993-09-13
PRIOR APPLICATION NUMBER: 07/955,243
PRIOR APPLICATION NUMBER: 07/955,243
PRIOR APPLICATION NUMBER: 07/955,243
PRIOR APPLICATION NUMBER: 07/955,243
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LENGTH: 788
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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Best Local (
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Publication No. US20030054554A1
PUBRAL INFORMATION:
PPLICANT: BECQUART, JEROME
TITLE OF INVENTION: ALBUMIN DERIVATIVES WITH THERAPEUTIC FUNCTIONS
FILE REFERENCE: 06832.1429-03
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Best Local Similarity
Matches 12; Conserv
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NUMBER OF SEQ ID NOS: 41
SOFTWARE: Patentin Ver. 2.1
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CURRENT FILING DATE: 2002-02-12
PRIOR APPLICATION NUMBER: 09/551,635
PRIOR FILING DATE: 2000-04-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: Protein fusion OTHER INFORMATION: prepro-HSA-VIV2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   203 DAHKSEVAHRFK 214
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FILING DATE: 1990-08-02
APPLICATION NUMBER: FR 89 10480
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TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 16:
                                                                                                                                                                                                                                                                                                                                                                                                                               12;
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; ORGANISM: Rattus norvegicus US-10-165-603-24
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; TYPE: PRT
; ORGANISM: Bovine Serum Albumin
US-10-045-170A-1
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                                                    SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 24
LENGTH: 608
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Qiu, Yongchang
APPLICANT: Qiu, Yongchang
APPLICANT: Wang, Jack
APPLICANT: Hewick, Rodney
TITLE OF INVENTION: ACID-LABILE ISOTOPE-CODED EXTRACTANT (ALICE) AND ITS USE IN QUANTI
TITLE OF INVENTION: MASS SPECTROMETRIC ANALYSIS OF PROTEIN MIXTURES
FILE REFERENCE: GIS412AUSA
CURRENT APPLICATION NUMBER: US/10/045,170A
CURRENT FILING DATE: 2001-10-22
CURRENT FILING DATE: 2001-10-22
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SEQ ID NO 89
LENGTH: 1184
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 24, Application US/10165603
Publication No. US20030021792A1
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Best Local Similarity
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                                                                                                                        PRIOR APPLICATION NUMBER: 60/297,021
PRIOR FILING DATE: 2001-06-08
PRIOR APPLICATION NUMBER: 60/305,117
PRIOR FILING DATE: 2001-07-12
NUMBER OF SEQ ID NOS: 33
                                                                                                                                                                                                                               APPLICANT: Stevens, Anthony C.
TITLE OF INVENTION: TISSUE-SPECIFIC ENDOTHELIAL MEMBRANE
TITLE OF INVENTION: PROTEINS
FILE REFERENCE: TOTECH.001A
CURRENT APPLICATION NUMBER: US/10/165,603
CURRENT FILING DATE: 2002-06-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/242643
PRIOR FILING DATE: 2000-10-23
NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Roben, Paul W. APPLICANT: Stevens, Anth
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                                           TYPE: PRT
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Pred. No. 0.0046;
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RESULT 49
US-10-076-071-4
; Sequence 4, Application US/10076071
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                                                                                                                                                                                                                            ; ORGANISM: Homo sapiens
US-09-845-726-1
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Publication No. US20030021792A1 
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Jackowski, George
APPLICANT: Jackowski, George
TITLE OF INVENTION: BIOPOLYMER MARKER INDICATIVE OF DISEASE STATE HAVING A MOLECULAR
RITLE OF INVENTION: OF 1424 DALTONS
FILE REFERENCE: 2132.033
CURRENT APPLICATION UNMBER: US/09/845,726
CURRENT APPLICATION WHEBER: 2001-04-30
RUMBER OF SEQ ID MOS: 1
SOFTWARE: PATENTIN VETBION 3.1
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 33
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 25
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CURRENT APPLICATION NUMBER: US/10/165,603
CURRENT FILING DATE: 2002-06-07
PRIOR APPLICATION NUMBER: 60/297,021
PRIOR FILING DATE: 2001-06-08
PRIOR APPLICATION NUMBER: 60/305,117
PRIOR FILING DATE: 2001-07-12
                                                                                                                                                                                            Query Match
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TITLE OF INVENTION: TISSUE-SPECIFIC ENDOTHELIAL MEMBRANE
TITLE OF INVENTION: PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Roben,
APPLICANT: Steven
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TYPE: PRT
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nes 10; Conserv
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100.0%; Pred. No. 7.8e-05;
Mismatches 0;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 59; DB 9; Pred. No. 0.0046;
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; TYPE: PRT ; ORGANISM: Homo sapiens US-10-076-071-4
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CURRENT FILING DATE: 2002-02-13
PRIOR APPLICATION NUMBER: 09/678,202
PRIOR FILING DATE: 2000-09-29
PRIOR APPLICATION NUMBER: 60/283,507
PRIOR APPLICATION NUMBER: 60/283,507
PRIOR APPLICATION NUMBER: 09/816,679
PRIOR FILING DATE: 2001-04-11
PRIOR FILING DATE: 2001-03-22
PRIOR FILING DATE: 1999-10-01
PRIOR APPLICATION NUMBER: 60/157,404
PRIOR APPLICATION NUMBER: 60/157,404
PRIOR APPLICATION NUMBER: 60/157,404
PRIOR APPLICATION NUMBER: 60/157,404
PRIOR FILING DATE: 1999-10-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Bar-Or, David
APPLICANT: Curtis, C. Ger
APPLICANT: Lau, Edward
APPLICANT: Rao, Nagaraja
APPLICANT: Winkley, James
APPLICANT: Winkley, James
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CURRENT FILING DATE: 2002-02-13
PRIOR APPLICATION NUMBER: 09/678,202
PRIOR FILING DATE: 2000-09-29
PRIOR APPLICATION NUMBER: 69/283,507
PRIOR APPLICATION NUMBER: 09/816,679
PRIOR APPLICATION NUMBER: 09/816,679
PRIOR APPLICATION NUMBER: 09/816,679
PRIOR FILING DATE: 2001-03-22
PRIOR FILING DATE: 2001-03-22
PRIOR FILING DATE: 2001-03-22
PRIOR APPLICATION NUMBER: 60/157,404
PRIOR FILLING DATE: 1999-10-01
PRIOR APPLICATION NUMBER: 60/157,404
PRIOR FILLING DATE: 1999-10-01
PRIOR FILLING DATE: 1999-10-01
PRIOR APPLICATION NUMBER: 60/211,078
PRIOR FILLING DATE: 2000-06-13
PRIOR APPLICATION NUMBER: 60/268,558
PRIOR FILLING DATE: 2001-02-13
NUMBER OF SEQ ID NOS: 9
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SEQ ID NO 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 5, Application US/10076071 Publication No. US20030060408A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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Best Local Similarity
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APPLICANT: Curtis, C. Ge
APPLICANT: Lau, Edward
APPLICANT: Rao, Nagaraje
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Lau, Edward
APPLICANT: Rao, Nagaraja K.R.
APPLICANT: Winkler, James V.
APPLICANT: Crook, Wannell M.
TITLE OF INVENTION: Metal-Binding Compounds and Uses Therefor FILE REFERENCE: 4172-3-2
CIPPERNT REFERENCE: 4172-3-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/211,078
PRIOR FILING DATE: 2000-06-13
PRIOR APPLICATION NUMBER: 60/268,558
PRIOR FILING DATE: 2001-02-13
NUMBER OF SEQ ID NOS: 9
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APPLICANT: Winkler, James V.
APPLICANT: Crook, Wannell M.
TITLE OF INVENTION: Metal-Binding Compounds and Uses Therefor
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100.0%; Pred. No. 7.8e-05;
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Search completed: April 11, 2003, 15:20:27 Job time : 16 secs
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; SEQ ID NO 5
; LENGTH: 10
; TYPE: PRT
; ORGANISM: Homo Bapiens
US-10-076-071-5
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                                                                                                                                Query Match 84.4%; Score 54; DB 9; Length 10; Best Local Similarity 100.0%; Pred. No. 0.00039; Matches 10; Conservative 0; Mismatches 0; Indels
                                                                   3 HKSEVAHRFK 12
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1 HKSEVAHRFK 10
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